(11) EP 1 074 617 A2

(12)

EUROPEAN PATENT APPLICATION

(43) Date of publication: 07.02.2001 Bulletin 2001/06

(21) Application number: 00116126.4

(51) Int.

C12N 15/12, C12N 15/11,

CI.7: C07K 14/47, C07K 16/18,

C12Q 1/68

(22) Date of filing: 28.07.2000

(84) Designated Contracting States: AT BE CH CY DE DK ES FI FR GB GR IE IT LI LU MC NL PT SE Designated Extension States: AL LT LV MK RO SI

(30) Priority: 29.07.1999 JP 24803699 27.08.1999 JP 30025399 11.01.2000 JP 2000118776 02.05.2000 JP 2000183767 09.06.2000 JP 2000241899

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(54) Primers for synthesising full-length cDNA and their use

(57)Primers for synthesizing fulllength cDNAs and their use are provided.

5602 cDNA encoding a human protein has been isolated and nucleotide sequences of 5'-, and 3' -ends of the cDNA have been determined. Furthermore, primers

synthesizing the full-length cDNA have been provided to clarify the function of the protein encoded by the cDNA. The full-length cDNA of the present invention containing the translation start site provides information useful for analyzing the functions of the protein.

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Description

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FIELD OF THE INVENTION

The present invention relates to a polynucleotide encoding a novel protein, a protein encoded by the polynucleotide, and new uses of these.

10 BACKGROUND OF THE INVENTION

Currently, the sequencing projects, the determination and analysis of the genomic DNA of various living organisms have been in progress all over the world. The whole genomic sequences of more than 10 species of prokaryotes, a lower eukaryote, yeast, and a multicellular eukaryote, C. elegans are already determined. As to human genome, which is supposed to be composed of three thousand million base pairs, the world wide cooperative projects have been under way to analyze it, and the whole structure is predicted to be determined by the years 2002-2003. The aim of the determination of genomic sequence is to reveal the functions of all genes and their regulation and to understand living organisms as a network of interactions between genes, proteins, cells or individuals through deducing the information in a genome, which is a blueprint of the highly complicated living organisms. To understand living organisms by utilizing the genomic information from various species is not only important as an academie subject, but also socially significant from the viewpoint of industrial application.

However, determination of genomic sequences itself cannot identify the functions of all genes. For example, as for yeast, only the function of approximately half of the 6000 genes, which is predicted based on the genomic sequence, was able to be deduced. As for human, the number of the genes is predicted to be approximately one hundred thousand. Therefore, it is desirable to establish "a high throughput analysis system of the gene functions" which allows us to identify rapidly and efficiently the functions of vast amounts of the genes obtained by the genomic sequencing.

Many genes in the eukaryotic genome are split by introns into multiple exons. Thus, it is difficult to predict correctly the structure of encoded protein solely based on genomic information. In contrast, cDNA, which is produced from mRNA that lacks introns, encodes a protein as a single continuous amino acid sequence and allows us to identify the primary structure of the protein easily. In human cDNA research, to date, more than one million ESTs (Expression Sequence Tags) are publicly available, and the ESTs presumably cover not less than 80% of all human genes.

The information of ESTs is utilized for analyzing the structure of human genome, or for predicting the exon-regions of genomic sequences or their expression profile. However, many human ESTs have been derived from proximal regions to the 3'-end of cDNA, and information around the 5'-end of mRNA is extremely little. Among these human cDNAs, the number of the corresponding mRNAs whose encoding protein sequences are deduced is approximately 7000, and further, the number of full-length therein is only 5500. Thus, even including cDNA registered as EST, the percentage of human cDNA obtained so far is estimated to be 10-15% of all the genes.

It is possible to identify the transcription start site of mRNA on the genomic sequence based on the 5'-end sequence of a full-length cDNA, and to analyze factors involved in the stability

of mRNA that is contained in the cDNA, or in its regulation of expression at the translation stage. Also, since a full-length cDNA contains ATG, the translation start site, in the 5'-region, it can be translated into a protein in a correct frame. Therefore, it is possible to produce a large amount of the protein encoded by the cDNA or to analyze biological activity of the expressed protein by utilizing an appropriate expression system. Thus, analysis of a full-length cDNA provides valuable information which complements the information from genome sequencing. Also, full-length cDNA clones that can be expressed are extremely valuable in empirical analysis of gene function and in industrial application.

Therefore, if a novel human full-length cDNA is isolated, it can be used for developing medicines for diseases in which the gene is involved. The protein encoded by the gene can be used as a drug by itself. Thus, it has great significance to obtain a full-length cDNA encoding a novel human protein.

In particular, human secretory proteins or membrane proteins would be useful by itself as a medicine like tissue plasminogen activator (TPA), or as a target of medicines like membrane receptors. In addition, genes for signal transduction-associated proteins (protein kinases, etc.) , glycoprotein-associated proteins, transcription-associated proteins, etc. are genes whose relationships to human diseases have been elucidated. Moreover, genes for diseaseassociated proteins form a gene group rich in genes whose relationships to human diseases have been elucidated.

Therefore, it has great significance to isolate novel full-length cDNA clones of human, only few of which has been isolated. Especially, isolation of a novel cDNA clone encoding a secretory protein or membrane protein is desired since the protein itself would be useful as a medicine, and also the clones potentially include a gene associated with diseases. In addition, genes encoding proteins that are associated with signal transduction, glycoprotein, transcription, or diseases are expected to be useful as target molecules for therapy, or as medicines themselves. These genes form a gene group predicted to be strongly associated with diseases. Thus, identification of the full-length cDNA clones encoding those proteins has great significance.

<u>SUMMARY OF THE INVENTION</u>

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An objective of the present invention is to provide a polynucleotide encoding a novel protein, 40 a protein encoded by said polynucleotide, and novel usages of these.

The inventors have developed a method for efficiently cloning a human full-length cDNA that is predicted by the ATGpr etc. to be a full-length cDNA clone, from a full-length-enriched cDNA library that is synthesized by the oligo-capping method. Then, the inventors determined the nucleotide sequence of the obtained cDNA clones from both 5'- and 3'- ends.

Furthermore, the inventors analyzed the obtained clones by the BLAST search of the databases, SwissProt (http://www.ebi.ac.uk/ebi_docsSwissProt_db/swisshome.html), (http: UniGene (http://www.ncbi.nlm.nih.gov/web/GenBank), and //www.ncbi.nlm.nih.gov/UniGene).

The full-length cDNA clones of the present invention have high fullness ratio since these were obtained by the combination of (1) construction of a full-length-enriched cDNA library that is synthesized by the oligo-capping method, and (2) a system in which the full-length ratio is evaluated from the nucleotide sequence of the 5'-end (selection based on the ATGpr, previously

removed complete sequences to ESTs). However, the primer of the present invention enables to obtain full-length cDNA easily without any specialized methods as in the described method. Homology analysis in which the analysis is carried out against a not-full-length cDNA fragment to postulate the function of a protein encoded by said fragment, is being commonly performed. However, since such analysis is based on the information of the fragment, it is not clear as to whether this fragment corresponds to a part that is functionally important in the protein. In other words, the reliability of the homology analysis based on the information of a fragment is doubtful, as information related to the structure of the whole protein is not available. However, the homology analysis of the present invention is conducted based on the information of a full-length cDNA comprising the whole coding region of the cDNA, and therefore, the homology of various portions of the protein can be analyzed. Hence, the reliability of the homology analysis has been dramatically improved in the present invention.

The inventors completed the invention by finding that it is possible to synthesize a novel full-length cDNA by using the combination of a primer that is designed based on the nucleotide sequence of the 5'-ends of the selected full-length cDNA clones and any of an oligo-dT primer or a 3'-primer that is designed based on the nucleotide sequence of the 3'-ends of the selected clones.

Thus, the present invention relates to primers described below, a method for synthesizing a polynucleotide using the primers, and polynucleotides obtained by the method.

First, the present invention relates to

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(1) use of an oligonucleotide as a primer for synthesizing the polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs: 1-5547 and SEQ ID NOs: 16111-16164, or the complementary strand thereof, wherein said oligonucleotide is complementary to said polynucleotide or the complementary strand thereof and comprises at least 15 nucleotides;

(2) a primer set for synthesizing polynucleotides, the primer set comprising an oligo-dT primer and an oligonucleotide complementary to the complementary strand of the polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs: 1-5547 and SEQ ID NOs: 16111-16164, wherein said oligonucleotide comprises at least 15 nucleotides; and

(3) a primer set for synthesizing polynucleotides, the primer set comprising a combination of an oligonucleotide comprising a nucleotide sequence complementary to the complementary strand of the polynucleotide comprising a 5'-end nucleotide sequence and an oligonucleotide comprising a nucleotide sequence complementary to the polynucleotide comprising a 3'-end nucleotide sequence, wherein said oligonucleotides comprise at least 15 nucleotides and wherein said combination of 5'-end nucleotide sequence / 3'-end nucleotide sequence is selected from the combinations of 5'-end nucleotide sequence / 3'-end nucleotide sequence set forth in the SEQ ID NOs in Tables 1 and 2.

Tables 1 and 2 shows names of clones obtained in the examples described later, comprising the polynucleotide of the present invention (Table 1; 5547 clones, Table 2; 54 clones), names of

nucleotide sequences at the 5'-end and 3'-end of the full-length cDNA, and their corresponding SEQ ID NOs. A blank indicates that the 3'-end sequence corresponding to the 5'-end sequence has not been determined for the same clone.

The SEQ ID NO of a 5'-end sequence is shown on the right side of the name of the 5'-end sequence, and the SEQ ID NO of a 3'-end sequence is shown on the right side of the name of the 3'-end sequence.

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name of	name of	SEQ ID	name of	SEQ ID
clone	5'-end	of 5'-end	3'-end	of 3'-end
	sequence	sequence	sequence	sequence
HEMBA1000005	F-HEMBA100000	5 1	R-HEMBA100000	5 5548
HEMBA1000012	F-HEMBA100001	2 2		
HEMBA1000020	F-HEMBA100002	0 3		
HEMBA1000030	F-HEMBA100003	0 4	R-HEMBA100003	0 5549
HEMBA1000042	F-HEMBA100004	2 5	R-HEMBA100004	2 5550
HEMBA1000046	F-HEMBA100004	6 6	R-HEMBA100004	6 5551
HEMBA1000050	F-HEMBA100005	50 7	R-HEMBA100005	0 5552
HEMBA1000076	F-HEMBA100007	6 8	R-HEMBA100007	6 5553
HEMBA1000111	F-HEMBA100011	1 9	R-HEMBA100011	1 5554
HEMBA1000129	F-HEMBA100012	9 10	R-HEMBA100012	9 5555
HEMBA1000141	F-HEMBA100014	11 11	R-HEMBA100014	1 5556
HEMBA1000150	F-HEMBA100015	50 12	R-HEMBA100015	0 5557
HEMBA1000156	F-HEMBA100015	6 13	R-nnnnnnnnnnn	n 5558
HEMBA1000158	F-HEMBA100015	8 14	R-HEMBA100015	8 5559
HEMBA1000168	F-HEMBA100016	8 15	R-nnnnnnnnnnn	n 5560

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		EP 107	4011	MZ		
	HEMBA1000180	F-HEMBA1000180	16	R-HEMBA1000180	5561	
	HEMBA1000185	F-HEMBA1000185	17	R-HEMBA1000185	5562	
	HEMBA1000193	F-HEMBA1000193	18	R-HEMBA1000193	5563	
5	HEMBA1000201	F-HEMBA1000201	19	R-HEMBA1000201	5564	
	HEMBA1000213	F-HEMBA1000213	20	R-HEMBA1000213	5565	
	HEMBA1000216	F-HEMBA1000216	21	R-HEMBA1000216	5566	
	HEMBA1000227	F-HEMBA1000227	22	R-nnnnnnnnnnn	5567	
10	HEMBA1000231	F-HEMBA1000231	23	R-HEMBA1000231	5568	
	HEMBA1000243	F-HEMBA1000243	24	R-HEMBA1000243	5569	
	HEMBA1000244	F-HEMBA1000244	25	R-HEMBA1000244	5570	
	HEMBA1000251	F-HEMBA1000251	26	R-HEMBA1000251	5571	
15	HEMBA1000264	F-HEMBA1000264	27	R-HEMBA1000264	5572	
	HEMBA1000280	F-HEMBA1000280	28	R-nnnnnnnnnnn	5573	
	HEMBA1000282	F-HEMBA1000282	29	R-HEMBA1000282	5574	
	HEMBA1000288	F-HEMBA1000288	30	R-HEMBA1000288	5575	
20	HEMBA1000290	F-HEMBA1000290	31	R-HEMBA1000290	5576	
	HEMBA1000302	F-HEMBA1000302	32	R-HEMBA1000302	5577	
	HEMBA1000303	F-HEMBA1000303	33	R-nnnnnnnnnnn	5578	
	HEMBA1000304	F-HEMBA1000304	34	R-nannananana	5579	
25	HEMBA1000307	F-HEMBA1000307	35	R-HEMBA1000307	5580	ĺ
	HEMBA1000327	F-HEMBA1000327	36			
	HEMBA1000333	F-HEMBA1000333	37	R-nnnnnnnnnnn	5581	
	HEMBA1000338	F-HEMBA1000338	38	R-HEMBA1000338	5582	
30	HEMBA1000351	F-HEMBA1000351	39	R-HEMBA1000351	5583	
	HEMBA1000355	F-HEMBA1000355	40	R-HEMBA1000355	5584	
	HEMBA1000356	F-HEMBA1000356	41			
	HEMBA1000357	F-HEMBA1000357	42	R-HEMBA1000357	5585	
35	HEMBA1000366	F-HEMBA1000366	43	R-HEMBA1000366	5586	
	HEMBA1000369	F-HEMBA1000369	44	R-HEMBA1000369	5587	
	HEMBA1000376	F-HEMBA1000376	45	R-HEMBA1000376	5588	
	HEMBA1000387	F-HEMBA1000387	46	R-HEMBA1000387	5589	
40	HEMBA1000390	F-HEMBA1000390	47	R-HEMBA1000390	5590	
•	HEMBA1000392	F-HEMBA1000392	48	R-HEMBA1000392	5591	
	HEMBA1000396	F-HEMBA1000396	49	R-HEMBA1000396	5592	
	HEMBA1000411	F-HEMBA1000411	50	R-HEMBA1000411	5593	
45	HEMBA1000418	F-HEMBA1000418	51	R-HEMBA1000418	5594	
	HEMBA1000422	F-HEMBA1000422	52	R-HEMBA1000422	5595	
	HEMBA1000428	F-HEMBA1000428	53	R-HEMBA1000428	5596	
	HEMBA1000434	F-HEMBA1000434	54	R-HEMBA1000434	5597	ļ
50	HEMBA1000442	F-HEMBA1000442	55	R-HEMBA1000442	5598	
	HEMBA1000456	F-HEMBA1000456	56	R-HEMBA1000456	5599	1
	HEMBA1000459	F-HEMBA1000459	57	R-HEMBA1000459	5600	-
	HEMBA1000460	F-HEMBA1000460	58	R-HEMBA1000460	5601	
55	HEMBA1000464	F-HEMBA1000464	59	R-HEMBA1000464	5602	-
	HEMBA1000469	F-HEMBA1000469	60	R-HEMBA1000469	5603	
	HEMBA1000488	F-HEMBA1000488	61	R-HEMBA1000488	5604	
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	HEMBA1000490	F-HEMBA1000490	62	R-HEMBA1000490	5605
	HEMBA1000491	F-HEMBA1000491	63	R-HEMBA1000491	5606
	HEMBA1000501	F-HEMBA1000501	64		
5	HEMBA1000504	F-HEMBA1000504	65	R-HEMBA1000504	5607
	HEMBA1000505	F-HEMBA1000505	66	R-HEMBA1000505	5608
	HEMBA1000508	F-HEMBA1000508	67	R-HEMBA1000508	5609
	HEMBA1000518	F-HEMBA1000518	68	R-HEMBA1000518	5610
10	HEMBA1000519	F-HEMBA1000519	69	R-HEMBA1000519	5611
	HEMBA1000520	F-HEMBA1000520	70	R-HEMBA1000520	5612
	HEMBA1000523	F-HEMBA1000523	71	R-HEMBA1000523	5613
	HEMBA1000531	F-HEMBA1000531	72	R-HEMBA1000531	5614
15	HEMBA1000534	F-HEMBA1000534	73	••	
	HEMBA1000540	F-HEMBA1000540	74	R-HEMBA1000540	5615
	HEMBA1000542	F-HEMBA1000542,	75		
	HEMBA1000545	F-HEMBA1000545	76	R-HEMBA1000545	5616
20	HEMBA1000555	F-HEMBA1000555	77	R-nnnnnnnnnn	5617
20	HEMBA1000557	F-HEMBA1000557	78	R-HEMBA1000557	5618
	HEMBA1000561	F-HEMBA1000561	79	R-HEMBA1000561	5619
	HEMBA1000563	F-HEMBA1000563	80	R-HEMBA1000563	5620
25	HEMBA1000568	F-HEMBA1000568	81	R-HEMBA1000568	5621
20	HEMBA1000569	F-HEMBA1000569	82	R-nnnnnnnnnnn	5622
	HEMBA1000575	F-HEMBA1000575	83	R-HEMBA1000575	5623
	HEMBA1000588	F-HEMBA1000588	84	R-HEMBA1000588	5624
30	HEMBA1000591	F-HEMBA1000591	85	R-HEMBA1000591	5625
30	HEMBA1000592	F-HEMBA1000592	86	R-HEMBA1000592	5626
	HEMBA1000594	F-HEMBA1000594	87	R-HEMBA1000594	5627
	HEMBA1000604	F-HEMBA1000604	88	R-HEMBA1000604	5628
25	HEMBA1000608	F-HEMBA1000608	89	R-HEMBA1000608	5629
35	HEMBA1000622	F-HEMBA1000622	90	R-HEMBA1000622	5630
	HEMBA1000636	F-HEMBA1000636	91	R-HEMBA1000636	5631
	HEMBA1000637	F-HEMBA1000637	92	R-HEMBA1000637	5632
40	HEMBA1000655	F-HEMBA1000655	93	R-HEMBA1000655	5633
40	HEMBA1000657	F-HEMBA1000657	94	R-HEMBA1000657	5634
	HEMBA1000662	F-HEMBA1000662	95	R-HEMBA1000662	5635
	HEMBA1000673	F-HEMBA1000673	96	R-HEMBA1000673	5636
	HEMBA1000682	F-HEMBA1000682	97	R-HEMBA1000682	5637
45	HEMBA1000686	F-HEMBA1000686	98	R-HEMBA1000686	5638
	HEMBA1000702	F-HEMBA1000702	99	R-HEMBA1000702	5639
	HEMBA1000705	F-HEMBA1000705	100	R-HEMBA1000705	5640
	HEMBA1000719	F-HEMBA1000719	101	R-HEMBA1000719	5641
50	HEMBA1000722	F-HEMBA1000722	102	R-HEMBA1000722	5642
	HEMBA1000726	F-HEMBA1000726	103	R-HEMBA1000726	5643
	HEMBA1000727	F-HEMBA1000727	104	R-HEMBA1000727	5644
	HEMBA1000747	F-HEMBA1000747	105	R-HEMBA1000747	5645
55	HEMBA1000749	F-HEMBA1000749	106	R-HEMBA1000749	5646
	HEMBA1000752	F-HEMBA1000752	107	R-HEMBA1000752	5647

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	HEMBA1000769	F-HEMBA1000769	108	R-HEMBA1000769	5648
	HEMBA1000773	F-HEMBA1000773	109	R-HEMBA1000773	5649
	HEMBA1000774	F-HEMBA1000774	110	R-HEMBA1000774	5650
5	HEMBA1000791	F-HEMBA1000791	111	R-HEMBA1000791	5651
	HEMBA1000817	F-HEMBA1000817	112	R-HEMBA1000817	5652
	HEMBA1000822	F-HEMBA1000822	113	R-HEMBA1000822	5653
	HEMBA1000827	F-HEMBA1000827	114	R-HEMBA1000827	5654
10	HEMBA1000843	F-HEMBA1000843	115	R-HEMBA1000843	5655
	HEMBA1000851	F-HEMBA1000851	116	R-HEMBA1000851	5656
	HEMBA1000852	F-HEMBA1000852	117	R-HEMBA1000852	5657
	HEMBA1000867	F-HEMBA1000867	118	R-HEMBA1000867	5658
15	HEMBA1000869	F-HEMBA1000869	119	R-HEMBA1000869	5659
10	HEMBA1000870	F-HEMBA1000870	120	R-HEMBA1000870	5660
	HEMBA1000870	F-HEMBA1000872	121		
	HEMBA1000872	F-HEMBA1000872,		R-HEMBA1000872	5661
			122	R-HEMBA1000876	5662
20	HEMBA1000908	F-HEMBA1000908	123	R-HEMBA1000908	5663
	HEMBA1000910	F-HEMBA1000910	124	R-HEMBA1000910	5664
	HEMBA1000918	F-HEMBA1000918	125	R-HEMBA1000918	5665
	HEMBA1000919	F-HEMBA1000919	126	R-HEMBA1000919	5666
25	HEMBA1000934	F-HEMBA1000934	127	R-HEMBA1000934	5667
	HEMBA1000942	F-HEMBA1000942	128	R-HEMBA1000942	5668
	HEMBA1000943	F-HEMBA1000943	129	R-HEMBA1000943	5669
	HEMBA1000946	F-HEMBA1000946	130	R-HEMBA1000946	5670
30	HEMBA1000960	F-HEMBA1000960	131	R-HEMBA1000960	5671
	HEMBA1000968	F-HEMBA1000968	132	R-HEMBA1000968	5672
	HEMBA1000971	F-HEMBA1000971	133	R-HEMBA1000971	5673
	HEMBA1000972	F-HEMBA1000972	134	R-HEMBA1000972	5674
35	HEMBA1000974	F-HEMBA1000974	135	R-HEMBA1000974	5675
	HEMBA1000975	F-HEMBA1000975	136	R-HEMBA1000975	5676
	HEMBA1000985	F-HEMBA1000985	137	R-HEMBA1000985	5677
	HEMBA1000986	F-HEMBA1000986	138	R-HEMBA1000986	5678
40	HEMBA1000991	F-HEMBA1000991	139	R-HEMBA1000991	5679
	HEMBA1001007	F-HEMBA1001007	140	R-HEMBA1001007	5680
	HEMBA1001008	F-HEMBA1001008	141	R-HEMBA1001008	5681
	HEMBA1001009	F-HEMBA1001009	142	R-HEMBA1001009	5682
45	HEMBA1001017	F-HEMBA1001017	143	R-HEMBA1001017	5683
	HEMBA1001019	F-HEMBA1001019	144	R-HEMBA1001019	5684
	HEMBA1001020	F-HEMBA1001020	145	R-HEMBA1001020	5685
	HEMBA1001022	F-HEMBA1001022	146	R-HEMBA1001022	5686
50	HEMBA1001024	F-HEMBA1001024	147	R-HEMBA1001024	5687
50	HEMBA1001026	F-HEMBA1001026	148	R-HEMBA1001026	5688
	HEMBA1001043	F-HEMBA1001043	149	R-nnnnnnnnnnn	5689
	HEMBA1001051	F-HEMBA1001051	150	R-HEMBA1001051	5690
55	HEMBA1001052	F-HEMBA1001052	151	R-HEMBA1001052	5691
55	HEMBA1001059	F-HEMBA1001059	152		
	HEMBA1001060	F-HEMBA1001060	153	R-HEMBA1001060	5692

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	·HEMBA1001071	F-HEMBA1001071	154	R-HEMBA1001071	5693
	HEMBA1001077	F-HEMBA1001077	155	R-HEMBA1001077	5694
	HEMBA1001080	F-HEMBA1001080	156	R-HEMBA1001080	5695
5	HEMBA1001085	F-HEMBA1001085	157	R-HEMBA1001085	5696
	HEMBA1001088	F-HEMBA1001088	158	R-HEMBA1001088	5697
	HEMBA1001094	F-HEMBA1001094	159	R-HEMBA1001094	5698
	HEMBA1001099	F-HEMBA1001099	160	R-HEMBA1001099	5699
10	HEMBA1001109	F-HEMBA1001109	161	R-HEMBA1001109	5700
	HEMBA1001121	F-HEMBA1001121	162	R-HEMBA1001121	5701
	HEMBA1001122	F-HEMBA1001122	163	R-HEMBA1001122	5702
	HEMBA1001123	F-HEMBA1001123	164	R-HEMBA1001123	5703
15	HEMBA1001133	F-HEMBA1001133	165	R-HEMBA1001133	5704
	HEMBA1001137	F-HEMBA1001137	166	R-HEMBA1001137	5705
	HEMBA1001140	F-HEMBA1001140,		R-HEMBA1001140	5706
	HEMBA1001172	F-HEMBA1001172	168	R-HEMBA1001172	5707
20	HEMBA1001174	F-HEMBA1001174	169	R-HEMBA1001174	5708
20	HEMBA1001197	F-HEMBA1001197	170	R-HEMBA1001197	5709
	HEMBA1001208	F-HEMBA1001208	171	R-HEMBA1001208	5710
	HEMBA1001213	F-HEMBA1001213	172	R INDINITIONS	0.10
25	HEMBA1001226	F-HEMBA1001226	173	R-HEMBA1001226	5711
25	HEMBA1001235	F-HEMBA1001235	174	R-HEMBA1001235	5712
	HEMBA1001247	F-HEMBA1001247	175	R-HEMBA1001247	5713
	HEMBA1001257	F-HEMBA1001257	176	R-HEMBA1001257	5714
20	HEMBA1001265	F-HEMBA1001265	177	R-HEMBA1001265	5715
30	HEMBA1001281	F-HEMBA1001281	178	R-nnnnnnnnnnn	5716
	HEMBA1001286	F-HEMBA1001286	179	R-HEMBA1001286	5717
	HEMBA1001289	F-HEMBA1001289	180	R-HEMBA1001289	5718
25	HEMBA1001294	F-HEMBA1001294	181	R-HEMBA1001294	5719
35	HEMBA1001299	F-HEMBA1001299	182	R-HEMBA1001299	5720
	HEMBA1001302	F-HEMBA1001302	183	R-HEMBA1001302	5721
	HEMBA1001303	F-HEMBA1001303	184	R-HEMBA1001303	5722
	HEMBA1001310	F-HEMBA1001310	185	R-HEMBA1001310	5723
40	HEMBA1001319	F-HEMBA1001319	186	R-HEMBA1001319	5724
	HEMBA1001323	F-HEMBA1001323	187	R-HEMBA1001323	5725
	HEMBA1001326	F-HEMBA1001326	188	R-HEMBA1001326	5726
	HEMBA1001327	F-HEMBA1001327	189	R-HEMBA1001327	5727
45	HEMBA1001330	F-HEMBA1001330	190	R-HEMBA1001330	5728
	HEMBA1001351	F-HEMBA1001351	191	R-HEMBA1001351	5729
	HEMBA1001361	F-HEMBA1001361	192	R-HEMBA1001361	5730
	HEMBA1001375	F-HEMBA1001375	193	R-HEMBA1001375	5731
50	HEMBA1001377	F-HEMBA1001377	194	R-HEMBA1001377	5732
	HEMBA1001383	F-HEMBA1001383	195	R-HEMBA1001383	5733
	HEMBA1001387	F-HEMBA1001387	196	R-HEMBA1001387	5734
	HEMBA1001388	F-HEMBA1001388	197	R-HEMBA1001388	5735
55	HEMBA1001391	F-HEMBA1001391	198	R-HEMBA1001391	5736
	HEMBA1001398	F-HEMBA1001398	199	R-HEMBA1001398	5737

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	HEMBA1001405	F-HEMBA1001405	200	R-HEMBA1001405	5738
	HEMBA1001407	F-HEMBA1001407	201	R-HEMBA1001407	5739
	HEMBA1001411	F-HEMBA1001411	202	R-HEMBA1001411	5740
5	HEMBA1001413	F-HEMBA1001413	203	R-HEMBA1001413	5741
	HEMBA1001415	F-HEMBA1001415	204	R-HEMBA1001415	5742
	HEMBA1001432	F-HEMBA1001432	205	R-HEMBA1001432	5743
	HEMBA1001433	F-HEMBA1001433	206	R-HEMBA1001433	5744
10	HEMBA1001435	F-HEMBA1001435	207	R-HEMBA1001435	5745
	HEMBA1001442	F-HEMBA1001442	208	R-HEMBA1001442	5746
	HEMBA1001446	F-HEMBA1001446	209	R-HEMBA1001446	5747
	HEMBA1001450	F-HEMBA1001450	210	R-HEMBA1001450	5748
15	HEMBA1001454	F-HEMBA1001454	21:1	R-HEMBA1001454	5749
	HEMBA1001455	F-HEMBA1001455	212	R-HEMBA1001455	5750
	HEMBA1001463	F-HEMBA1001463,	213	R-HEMBA1001463	5751
	HEMBA1001476	F-HEMBA1001476	214	R-HEMBA1001476	5752
20	HEMBA1001478	F-HEMBA1001478	215	R-HEMBA1001478	5753
	HEMBA1001497	F-HEMBA1001497	216	R-HEMBA1001497	5754
	HEMBA1001510	F-HEMBA1001510	217	R-HEMBA1001510	5755
	HEMBA1001515	F-HEMBA1001515	218	R-HEMBA1001515	5756
25	HEMBA1001517	F-HEMBA1001517	219	R-HEMBA1001517	5757
20	HEMBA1001522	F-HEMBA1001522	220	R-HEMBA1001522	5758
	HEMBA1001526	F-HEMBA1001526	221	R-HEMBA1001526	5759
	HEMBA1001533	F-HEMBA1001533	222	R-HEMBA1001533	5760
30	HEMBA1001557	F-HEMBA1001557	223	R-HEMBA1001557	5761
30	HEMBA1001566	F-HEMBA1001566	224	R-HEMBA1001566	5762
	HEMBA1001569	F-HEMBA1001569	225	R-HEMBA1001569	5763
	HEMBA1001570	F-HEMBA1001570	226	R-HEMBA1001570	5764
05	HEMBA1001579	F-HEMBA1001579	227	R-HEMBA1001579	5 765
35	HEMBA1001581	F-HEMBA1001581	228	R-HEMBA1001581	5766
	HEMBA1001585	F-HEMBA1001585	229	R-HEMBA1001585	5767
	HEMBA1001589	F-HEMBA1001589	230	R-HEMBA1001589	5768
	HEMBA1001595	F-HEMBA1001595	231	R-HEMBA1001595	5769
40	HEMBA1001608	F-HEMBA1001608	232	R-HEMBA1001608	5770
	HEMBA1001620	F-HEMBA1001620	233	R-HEMBA1001620	5771
	HEMBA1001635	F-HEMBA1001635	234	R-nnnnnnnnnnn	5772
	HEMBA1001636	F-HEMBA1001636	235	R-HEMBA1001636	5773
45	HEMBA1001640	F-HEMBA1001640	236	R-HEMBA1001640	5774
	HEMBA1001647	F-HEMBA1001647	237		
	HEMBA1001651	F-HEMBA1001651	238	R-nnnnnnnnnnn	5775
	HEMBA1001655	F-HEMBA1001655	239	R-HEMBA1001655	5776
50	HEMBA1001658	F-HEMBA1001658	240	R-HEMBA1001658	5777
	HEMBA1001661	F-HEMBA1001661	241	R-HEMBA1001661	5778
	HEMBA1001672	F-HEMBA1001672	242	R-HEMBA1001672	5779
	HEMBA1001675	F-HEMBA1001675	243	R-HEMBA1001675	5780
55	HEMBA1001678	F-HEMBA1001678	244	R-HEMBA1001678	5781
	HEMBA1001681	F-HEMBA1001681	245	R-HEMBA1001681	5782

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	HEMBA1001702	F-HEMBA1001702	246	R-HEMBA1001702	5783
	HEMBA1001709	F-HEMBA1001709	247	R-HEMBA1001709	5784
	HEMBA1001711	F-HEMBA1001711	248	R-HEMBA1001711	5785
5	HEMBA1001712	F-HEMBA1001712	249	R-HEMBA1001712	5786
	HEMBA1001714	F-HEMBA1001714	250	R-HEMBA1001714	5787
	HEMBA1001718	F-HEMBA1001718	251	R-HEMBA1001718	5788
	HEMBA1001723	F-HEMBA1001723	252	R-HEMBA1001723	5789
10	HEMBA1001731	F-HEMBA1001731	253	R-HEMBA1001731	5790
	HEMBA1001734	F-HEMBA1001734	254	R-HEMBA1001734	5791
	HEMBA1001744	F-HEMBA1001744	255	R-HEMBA1001744	5792
	HEMBA1001745	F-HEMBA1001745	256	R-HEMBA1001745	5793
15	HEMBA1001746	F-HEMBA1001746	257	R-HEMBA1001746	5794
10	HEMBA1001761	F-HEMBA1001761	258	R-HEMBA1001761	5795
	HEMBA1001781	F-HEMBA1001781	259	R-HEMBA1001781	5796
	HEMBA1001784	F-HEMBA1001784	260	R-HEMBA1001784	5797
20	HEMBA1001791	F-HEMBA1001791	261	R-HEMBA1001791	5798
20	HEMBA1001800	F-HEMBA1001800	262	R-HEMBA1001800	5799
	HEMBA1001803	F-HEMBA1001803	263	R-HEMBA1001803	5800
	HEMBA1001804	F-HEMBA1001804	264	R-nnnnnnnnn	5801
	HEMBA1001804	F-HEMBA1001808	265	R-HEMBA1001808	5802
25	HEMBA1001809	F-HEMBA1001809	266	R-HEMBA1001809	5803
	HEMBA1001805	F-HEMBA1001815	267	R-HEMBA1001815	5804
	HEMBA1001819	F-HEMBA1001819	268	R-HEMBA1001819	5805
	HEMBA1001819	F-HEMBA1001820	269	R-HEMBA1001820	5806
30	HEMBA1001822	F-HEMBA1001822	270	R-nnnnnnnnnnnn	5807
	HEMBA1001824	F-HEMBA1001824	271	R-HEMBA1001824	5808
	HEMBA1001824	F-HEMBA1001835	272	R-HEMBA1001835	5809
	HEMBA1001844	F-HEMBA1001844	273	R-HEMBA1001844	5810
35	HEMBA1001847	F-HEMBA1001847	274	R-HEMBA1001847	5811
	HEMBA1001861	F-HEMBA1001861	275	R-HEMBA1001861	5812
	HEMBA1001864	F-HEMBA1001864	276	R-HEMBA1001864	5813
	HEMBA1001866	F-HEMBA1001866	277	R-HEMBA1001866	5814
40	HEMBA1001869	F-HEMBA1001869	278	R-nnnnnnnnnnn	5815
	HEMBA1001888	F-HEMBA1001888	279	R-HEMBA1001888	5816
	HEMBA1001896	F-HEMBA1001896	280	R-HEMBA1001896	5817
	HEMBA1001910	F-HEMBA1001910	281	R-HEMBA1001910	5818
45	HEMBA1001912	F-HEMBA1001912	282	R-HEMBA1001912	5819
	HEMBA1001913	F-HEMBA1001913	283	R-HEMBA1001913	5820
	HEMBA1001915	F-HEMBA1001915	284	R-HEMBA1001915	5821
	HEMBA1001918	F-HEMBA1001918	285	R-HEMBA1001918	5822
50	HEMBA1001921	F-HEMBA1001921	286	R-HEMBA1001921	5823
	HEMBA1001939	F-HEMBA1001939	287	R-HEMBA1001939	5824
	HEMBA1001940	F-HEMBA1001940	288	R-HEMBA1001940	5825
	HEMBA1001942	F-HEMBA1001942	289	R-HEMBA1001942	5826
55	HEMBA1001945	F-HEMBA1001945	290	R-HEMBA1001945	5827
	HEMBA1001950	F-HEMBA1001950	291	R-HEMBA1001950	5828
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	HEMBA1001960	F-HEMBA1001960	292	R-HEMBA1001960	5829
	HEMBA1001962	F-HEMBA1001962	293	R-HEMBA1001962	5830
	HEMBA1001964	F-HEMBA1001964	294	R-HEMBA1001964	5831
5	HEMBA1001967	F-HEMBA1001967	295	R-HEMBA1001967	5832
	HEMBA1001979	F-HEMBA1001979	296	R-HEMBA1001979	5833
	HEMBA1001987	F-HEMBA1001987	297	R-HEMBA1001987	5834
	HEMBA1001991	F-HEMBA1001991	298	R-HEMBA1001991	5835
10	HEMBA1002003	F-HEMBA1002003	299	R-HEMBA1002003	5836
70	HEMBA1002008	F-HEMBA1002008	300	R-HEMBA1002008	5837
	HEMBA1002018	F-HEMBA1002018	301	R-HEMBA1002018	5838
	HEMBA1002022	F-HEMBA1002022	302	R-HEMBA1002022	5839
15	HEMBA1002035	F-HEMBA1002035	303	R-HEMBA1002035	5840
15	HEMBA1002039	F-HEMBA1002039	304	R-HEMBA1002039	5841
	HEMBA1002049	F-HEMBA1002049	305	R-HEMBA1002049	5842
	HEMBA1002084	F-HEMBA1002084	306	R-HEMBA1002084	5843
••	HEMBA1002092	F-HEMBA1002092	307	R-HEMBA1002092	5844
20	HEMBA1002100	F-HEMBA1002100	308	R-HEMBA1002100	5845
	HEMBA1002102	F-HEMBA1002102	309	R-HEMBA1002102	5846
	HEMBA1002113	F-HEMBA1002113	310	R-HEMBA1002113	5847
	HEMBA1002119	F-HEMBA1002119	311	R-HEMBA1002119	5848
25	HEMBA1002125	F-HEMBA1002125	312	R-HEMBA1002125	5849
	HEMBA1002139	F-HEMBA1002139	313	R-HEMBA1002139	5850
	HEMBA1002144	F-HEMBA1002144	314	R-HEMBA1002144	5851
	HEMBA1002150	F-HEMBA1002150	315	R-HEMBA1002150	5852
30	HEMBA1002151	F-HEMBA1002151	316	R-HEMBA1002151	5853
	HEMBA1002153	F-HEMBA1002153	317	R-HEMBA1002153	5854
	HEMBA1002160	F-HEMBA1002160	318	R-HEMBA1002160	5855
	HEMBA1002161	F-HEMBA1002161	319	R-HEMBA1002161	5856
35	HEMBA1002162	F-HEMBA1002162	320	R-HEMBA1002162	5857
	HEMBA1002166	F-HEMBA1002166	321	R-HEMBA1002166	5858
	HEMBA1002177	F-HEMBA1002177	322	R-HEMBA1002177	5859
	HEMBA1002185	F-HEMBA1002185	323	R-HEMBA1002185	5860
40	HEMBA1002189	F-HEMBA1002189	324	R-HEMBA1002189	5861
	HEMBA1002191	F-HEMBA1002191	325	R-HEMBA1002191	5862
	HEMBA1002199	F-HEMBA1002199	326	R-HEMBA1002199	5863
	HEMBA1002204	F-HEMBA1002204	327	R-HEMBA1002204	5864
45	HEMBA1002212	F-HEMBA1002212	328	R-HEMBA1002212	5865
	HEMBA1002215	F-HEMBA1002215	329	R-HEMBA1002215	5866
	HEMBA1002226	F-HEMBA1002226	330	R-HEMBA1002226	5867
	HEMBA1002229	F-HEMBA1002229	331	R-HEMBA1002229	5868
50	HEMBA1002237	F-HEMBA1002237	332	R-HEMBA1002237	5869
	HEMBA1002241	F-HEMBA1002241	333	D 11D10011000000	5070
	HEMBA1002253	F-HEMBA1002253	334	R-HEMBA1002253	5870
	HEMBA1002257	F-HEMBA1002257	335	R-HEMBA1002257	5871
55	HEMBA1002265	F-HEMBA1002265	336	D. HEMDA 10000CT	E070
	HEMBA1002267	F-HEMBA1002267	337	R-HEMBA1002267	5872

	HEMRA1009970	F-HEMBA1002270	338	R-HEMBA1002270	5873
					5874
	HEMBA1002321	F-HEMBA1002321	339	R-HEMBA1002321	
_		F-HEMBA1002328	340	R-HEMBA1002328	5875
5	HEMBA1002337	F-HEMBA1002337	341	R-HEMBA1002337	5876
	HEMBA1002341	F-HEMBA1002341	342	R-HEMBA1002341	5877
	HEMBA1002348	F-HEMBA1002348	343	R-HEMBA1002348	5878
	HEMBA1002349	F-HEMBA1002349	344	R-HEMBA1002349	5879
10	HEMBA1002363	F-HEMBA1002363	345	R-nnnnnnnnnnn	5880
	HEMBA1002381	F-HEMBA1002381	346	R-HEMBA1002381	5881
	HEMBA1002389	F-HEMBA1002389	347	R-HEMBA1002389	5882
	HEMBA1002417	F-HEMBA1002417	348	R-HEMBA1002417	5883
15	HEMBA1002419	F-HEMBA1002419	349	R-HEMBA1002419	5884
	HEMBA1002430	F-HEMBA1002430	350	R-HEMBA1002430	5885
	HEMBA1002439	F-HEMBA1002439,	351	R-HEMBA1002439	5886
	HEMBA1002458	F-HEMBA1002458	352	R-HEMBA1002458	5887
20	HEMBA1002460	F-HEMBA1002460	353	R-HEMBA1002460	5888
20	HEMBA1002462	F-HEMBA1002462	354	R-HEMBA1002462	5889
	HEMBA1002469	F-HEMBA1002469	355		
	HEMBA1002475	F-HEMBA1002475	356	R-nnnnnnnnnnn	5890
٥٢	HEMBA1002477	F-HEMBA1002477	357	R-HEMBA1002477	5891
25	HEMBA1002486	F-HEMBA1002486	358	R-HEMBA1002486	5892
	HEMBA1002495	F-HEMBA1002495	359	R-HEMBA1002495	5893
	HEMBA1002498	F-HEMBA1002498	360	R-HEMBA1002498	5894
	HEMBA1002503	F-HEMBA1002503	361	R-HEMBA1002503	5895
30	HEMBA1002508	F-HEMBA1002508	362	R-HEMBA1002508	5896
	HEMBA1002513	F-HEMBA1002513	363	R-nnnnnnnnnnn	5897
	HEMBA1002515	F-HEMBA1002515	364	R-HEMBA1002515	5898
	HEMBA1002538	F-HEMBA1002538	365	R-HEMBA1002538	5899
35	HEMBA1002542	F-HEMBA1002542	366	R-HEMBA1002542	5900
	HEMBA1002547	F-HEMBA1002547	367	R-HEMBA1002547	5901
	HEMBA1002552	F-HEMBA1002552	368	R-HEMBA1002552	5902
	HEMBA1002555	F-HEMBA1002555	369	R-HEMBA1002555	5903
40	HEMBA1002558	F-HEMBA1002558	370	R-HEMBA1002558	5904
	HEMBA1002561	F-HEMBA1002561	371	R-HEMBA1002561	5905
	HEMBA1002569	F-HEMBA1002569	372	R-nnnnnnnnnnn	5906
	HEMBA1002583	F-HEMBA1002583	373	R-HEMBA1002583	5907
45	HEMBA1002590	F-HEMBA1002590	374	R-HEMBA1002590	5908
	HEMBA1002592	F-HEMBA1002592	375	R-HEMBA1002592	5909
	HEMBA1002609	F-HEMBA1002609	376		
	HEMBA1002621	F-HEMBA1002621	377	R-HEMBA1002621	5910
50	HEMBA1002624	F-HEMBA1002624	378	R-HEMBA1002624	5911
30	HEMBA1002628	F-HEMBA1002628	379	R-HEMBA1002628	5912
	HEMBA1002629	F-HEMBA1002629	380	R-HEMBA1002629	5913
	HEMBA1002645	F-HEMBA1002645	381	R-HEMBA1002645	5914
	HEMBA1002651	F-HEMBA1002651	382	R-HEMBA1002651	5915
55	HEMBA1002659	F-HEMBA1002659	383	R-HEMBA1002659	5916

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	HEMBA1002661	F-HEMBA1002661	384	R-HEMBA1002661	5917
	HEMBA1002666	F-HEMBA1002666	385	R-HEMBA1002666	5918
	HEMBA1002678	F-HEMBA1002678	386	R-HEMBA1002678	5919
5	HEMBA1002679	F-HEMBA1002679	387	R-nnnnnnnnnnn	5920
	HEMBA1002688	F-HEMBA1002688	388	R-HEMBA1002688	5921
	HEMBA1002696	F-HEMBA1002696	389	R-HEMBA1002696	5922
	HEMBA1002703	F-HEMBA1002703	390		
10	HEMBA1002712	F-HEMBA1002712	391	R-HEMBA1002712	5923
	HEMBA1002716	F-HEMBA1002716	392	R-HEMBA1002716	5924
	HEMBA1002728	F-HEMBA1002728	393	R-HEMBA1002728	5925
	HEMBA1002730	F-HEMBA1002730	394	R-HEMBA1002730	5926
15	HEMBA1002742	F-HEMBA1002742	395	R-HEMBA1002742	5927
10	HEMBA1002746	F-HEMBA1002746	396	R-HEMBA1002746	5928
	HEMBA1002748	F-HEMBA1002748,	397	R-HEMBA1002748	5929
	HEMBA1002750	F-HEMBA1002750	398	R-HEMBA1002750	5930
00	HEMBA1002768	F-HEMBA1002768	399	R-HEMBA1002768	5931
20	HEMBA1002770	F-HEMBA1002770	400	R-HEMBA1002770	5932
	HEMBA1002777	F-HEMBA1002777	401	R-HEMBA1002777	5933
	HEMBA1002779	F-HEMBA1002779	402	R-HEMBA1002779	5934
	HEMBA1002780	F-HEMBA1002780	403	R-HEMBA1002780	593 5
25	HEMBA1002794	F-HEMBA1002794	404	R-HEMBA1002794	5936
	HEMBA1002801	F-HEMBA1002801	405	R-HEMBA1002801	5937
	HEMBA1002810	F-HEMBA1002810	406	R-HEMBA1002810	5938
	HEMBA1002816	F-HEMBA1002816	407	R-HEMBA1002816	5939
30	HEMBA1002818	F-HEMBA1002818	408	R ILMDRIVOZOIO	0000
	HEMBA1002826	F-HEMBA1002826	409	R-HEMBA1002826	5940
	HEMBA1002833	F-HEMBA1002833	410	R-HEMBA1002833	5941
	HEMBA1002850	F-HEMBA1002850	411	R-HEMBA1002850	5942
35	HEMBA1002863	F-HEMBA1002863	412	R-HEMBA1002863	5943
	HEMBA1002876	F-HEMBA1002876	413	R-HEMBA1002876	5944
	HEMBA1002886	F-HEMBA1002886	414	R-HEMBA1002886	5945
	HEMBA1002896	F-HEMBA1002896	415	R-HEMBA1002896	5946
40	HEMBA1002921	F-HEMBA1002921	416	R-HEMBA1002921	5947
	HEMBA1002924	F-HEMBA1002924	417	R-HEMBA1002924	5948
	HEMBA1002934	•	418	R-HEMBA1002934	5949
	HEMBA1002935	F-HEMBA1002935	419	R-HEMBA1002935	5950
45	HEMBA1002937	F-HEMBA1002937	420	R-HEMBA1002937	5951
	HEMBA1002939	F-HEMBA1002939	421	R-HEMBA1002939	5952
	HEMBA1002944	F-HEMBA1002944	422	R-HEMBA1002944	5953
	HEMBA1002951	F-HEMBA1002951	423	R-HEMBA1002951	5954
50	HEMBA1002954	F-HEMBA1002954	424	R-HEMBA1002954	5955
	HEMBA1002968	F-HEMBA1002968	425	R-HEMBA1002968	5956
	HEMBA1002970	F-HEMBA1002970	426	R-HEMBA1002970	5957
	HEMBA1002971	F-HEMBA1002971	427	R-HEMBA1002971	5958
66	HEMBA1002973	F-HEMBA1002973	428	R-HEMBA1002973	5959
55	HEMBA1002917	F-HEMBA1002997	429	R-nnnnnnnnnnn	5960
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	HEMBA1002999	F-HEMBA1002999	430	R-HEMBA1002999	5961	
	HEMBA1003021	F-HEMBA1003021	431	R-HEMBA1003021	5962	
	HEMBA1003033	F-HEMBA1003033	432	R-HEMBA1003033	5963	
5	HEMBA1003034	F-HEMBA1003034	433	R-HEMBA1003034	5964	
	HEMBA1003035	F-HEMBA1003035	434	R-HEMBA1003035	5965	
	HEMBA1003037	F-HEMBA1003037	435	R-HEMBA1003037	5966	
	HEMBA1003041	F-HEMBA1003041	436	R-HEMBA1003041	5967	
10	HEMBA1003046	F-HEMBA1003046	437	R-HEMBA1003046	5968	
	HEMBA1003064	F-HEMBA1003064	438	R-HEMBA1003064	5969	
	HEMBA1003067	F-HEMBA1003067	439	R-HEMBA1003067	5970	
	HEMBA1003071	F-HEMBA1003071	440	R-HEMBA1003071	5971	
15	HEMBA1003077	F-HEMBA1003077	441	R-HEMBA1003077	5972	
	HEMBA1003078	F-HEMBA1003078	442	R-HEMBA1003078	5973	
	HEMBA1003079	F-HEMBA1003079,	443	R-HEMBA1003079	5974	
	HEMBA1003083	F-HEMBA1003083	444	R-HEMBA1003083	5975	
20	HEMBA1003086	F-HEMBA1003086	445	R-HEMBA1003086	5976	
	HEMBA1003096	F-HEMBA1003096	446	R-HEMBA1003096	5977	
	HEMBA1003098	F-HEMBA1003098	447	R-HEMBA1003098	5978	
	HEMBA1003117	F-HEMBA1003117	448	R-HEMBA1003117	5979	
25	HEMBA1003129	F-HEMBA1003129	449	R-HEMBA1003129	5980	
	HEMBA1003133	F-HEMBA1003133	450	R-HEMBA1003133	5981	
	HEMBA1003136	F-HEMBA1003136	451	R-HEMBA1003136	5982	
	HEMBA1003142	F-HEMBA1003142	452	R-HEMBA1003142	5983	
30	HEMBA1003148	F-HEMBA1003148	453	R-HEMBA1003148	5984	
	HEMBA1003166	F-HEMBA1003166	454	R-HEMBA1003166	5985	
	HEMBA1003175	F-HEMBA1003175	455	R-HEMBA1003175	5986	
	HEMBA1003179	F-HEMBA1003179	456			
35	HEMBA1003197	F-HEMBA1003197	457	R-HEMBA1003197	5987	
	HEMBA1003199	F-HEMBA1003199	458	R-HEMBA1003199	5988	
	HEMBA1003202	F-HEMBA1003202	459	R-HEMBA1003202	5989	
	HEMBA1003204	F-HEMBA1003204	460	R-HEMBA1003204	5990	
40	HEMBA1003212	F-HEMBA1003212	461	R-HEMBA1003212	5991	
,,,	HEMBA1003220	F-HEMBA1003220	462	R-HEMBA1003220	5992	
	HEMBA1003222	F-HEMBA1003222	463	R-HEMBA1003222	5993	
	HEMBA1003229	F-HEMBA1003229	464	R-HEMBA1003229	5994	ļ
45	HEMBA1003235	F-HEMBA1003235	465	R-HEMBA1003235	5995	
40	HEMBA1003250	F-HEMBA1003250	466	R-HEMBA1003250	5996	
	HEMBA1003257	F-HEMBA1003257	467	R-HEMBA1003257	5997	
	HEMBA1003273	F-HEMBA1003273	468	R-HEMBA1003273	5998	
50	HEMBA1003276	F-HEMBA1003276	469	R-HEMBA1003276	5999	
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	HEMBA1003281	F-HEMBA1003281	471	R-HEMBA1003281	6001	
	HEMBA1003286	F-HEMBA1003286	472	D MEMBALAAAA	2000	
55	HEMBA1003291	F-HEMBA1003291	473	R-HEMBA1003291	6002	
55	HEMBA1003296	F-HEMBA1003296	474	R-HEMBA1003296	6003	1
	HEMBA1003304	F-HEMBA1003304	475	R-HEMBA1003304	6004	

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	HEMBA1003309	F-HEMBA1003309	476	R-HEMBA1003309	6005
	HEMBA1003314	F-HEMBA1003314	477	R-HEMBA1003314	6006
	HEMBA1003322	F-HEMBA1003322	478	R-HEMBA1003322	6007
5	HEMBA1003327	F-HEMBA1003327	479	R-HEMBA1003327	6008
	HEMBA1003328	F-HEMBA1003328	480	R-HEMBA1003328	6009
	HEMBA1003330	F-HEMBA1003330	481	R-HEMBA1003330	6010
	HEMBA1003348	F-HEMBA1003348	482	R-HEMBA1003348	6011
10	HEMBA1003369	F-HEMBA1003369	483	R-HEMBA1003369	6012
	HEMBA1003370	F-HEMBA1003370	484	R-HEMBA1003370	6013
	HEMBA1003373	F-HEMBA1003373	485	R-HEMBA1003373	6014
	HEMBA1003376	F-HEMBA1003376	486	R-HEMBA1003376	6015
45	HEMBA1003380	F-HEMBA1003380	487	R-HEMBA1003380	6016
15	HEMBA1003384	F-HEMBA1003384	488	R-HEMBA1003384	
	HEMBA1003395	F-HEMBA1003395,			6017
	HEMBA1003402	F-HEMBA1003402	489	R-HEMBA1003395	6018
	HEMBA1003402	F-HEMBA1003403	490	R-HEMBA1003402	6019
20	HEMBA1003408	F-HEMBA1003408	491		2002
	HEMBA1003417		492	R-nnnnnnnnnnnn	6020
	HEMBA1003417	F-HEMBA1003417	493	R-HEMBA1003417	6021
		F-HEMBA1003418	494	R-HEMBA1003418	6022
25	HEMBA1003433 HEMBA1003447	F-HEMBA1003433	495	R-HEMBA1003433	6023
	HEMBA1003447	F-HEMBA1003447 F-HEMBA1003461	496	D 10000401	C004
,	HEMBA1003463	F-HEMBA1003463	497	R-HEMBA1003461	6024
	HEMBA1003480	F-HEMBA1003480	498	R-HEMBA1003463	6025
30	HEMBA1003528	F-HEMBA1003528	499	R-HEMBA1003480	6026
	HEMBA1003531		500	R-HEMBA1003528	6027
	HEMBA1003531	F-HEMBA1003531	501	R-HEMBA1003531	6028
	HEMBA1003545	F-HEMBA1003538 F-HEMBA1003545	502	R-HEMBA1003538	6029
35	HEMBA1003548		503	R-HEMBA1003545	6030
	HEMBA1003555	F-HEMBA1003548 F-HEMBA1003555	504	R-HEMBA1003548	6031
	HEMBA1003556	F-HEMBA1003556	505	R-HEMBA1003555	6032
	HEMBA1003560		506	R-HEMBA1003556	6033
40	HEMBA1003568	F-HEMBA1003560	507	R-HEMBA1003560	6034
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45	HEMBA1003581	F-HEMBA1003579	511	R-HEMBA1003579	6038
45	HEMBA1003591	F-HEMBA1003581	512	R-HEMBA1003581	6039
	HEMBA1003595	F-HEMBA1003591	513	R-HEMBA1003591	6040
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		F-HEMBA1003597	515	R-HEMBA1003597	6042
50	HEMBA1003598	F-HEMBA1003598	516	R-HEMBA1003598	6043
	HEMBA1003615	F-HEMBA1003615	517	R-HEMBA1003615	6044
	HEMBA1003617	F-HEMBA1003617	518	R-HEMBA1003617	6045
	HEMBA1003621	F-HEMBA1003621	519	R-HEMBA1003621	6046
55	HEMBA1003622	F-HEMBA1003622	520	R-HEMBA1003622	6047
	HEMBA1003630	F-HEMBA1003630	521	R-HEMBA1003630	6048

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•	HEMBA1003637	F-HEMBA1003637	522	R-HEMBA1003637	6049
	HEMBA1003640	F-HEMBA1003640	523	R-HEMBA1003640	6050
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5	HEMBA1003646	F-HEMBA1003646	525	R-HEMBA1003646	6052
	HEMBA1003656	F-HEMBA1003656	526	R-HEMBA1003656	6053
	HEMBA1003662	F-HEMBA1003662	527	R-HEMBA1003662	6054
	HEMBA1003667	F-HEMBA1003667	528	R-HEMBA1003667	6055
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	HEMBA1003684	F-HEMBA1003684	531	R-HEMBA1003684	6058
	HEMBA1003690	F-HEMBA1003690	532	R-HEMBA1003690	6059
15	HEMBA1003692	F-HEMBA1003692	533	R-HEMBA1003692	6060
,,	HEMBA1003711	F-HEMBA1003711	534	R-HEMBA1003711	6061
	HEMBA1003714	F-HEMBA1003714,	535	R-HEMBA1003714	6062
	HEMBA1003715	F-HEMBA1003715	536	R-HEMBA1003715	6063
20	HEMBA1003720	F-HEMBA1003720	537	R-HEMBA1003720	6064
20	HEMBA1003725	F-HEMBA1003725	538	R-HEMBA1003725	6065
	HEMBA1003729	F-HEMBA1003729	539	R-HEMBA1003729	6066
	HEMBA1003733	F-HEMBA1003733	540	R-HEMBA1003733	6067
0.5	HEMBA1003742	F-HEMBA1003742	541	R-HEMBA1003742	6068
25	HEMBA1003758	F-HEMBA1003758	542	R-HEMBA1003758	6069
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	HEMBA1003773	F-HEMBA1003773	544	R-HEMBA1003773	6071
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	HEMBA1003803	F-HEMBA1003803	548	R-HEMBA1003803	6075
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35	HEMBA1003805	F-HEMBA1003805	550	R-HEMBA1003805	6077
	HEMBA1003807	F-HEMBA1003807	551	R-HEMBA1003807	6078
	HEMBA1003827	F-HEMBA1003827	552		
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40	HEMBA1003838	F-HEMBA1003838	554	R-HEMBA1003838	6080
	HEMBA1003856	F-HEMBA1003856	555	R-HEMBA1003856	6081
	HEMBA1003864	F-HEMBA1003864	556	R-HEMBA1003864	6082
	HEMBA1003866	F-HEMBA1003866	557	R-HEMBA1003866	6083
45	HEMBA1003879	F-HEMBA1003879	558	R-HEMBA1003879	6084
	HEMBA1003880	F-HEMBA1003880	559	R-HEMBA1003880	6085
	HEMBA1003885	F-HEMBA1003885	560	R-HEMBA1003885	6086
	HEMBA1003893	F-HEMBA1003893	561	R-HEMBA1003893	6087
50	HEMBA1003902	F-HEMBA1003902	562	R-HEMBA1003902	6088
	HEMBA1003908	F-HEMBA1003908	563	R-HEMBA1003908	6089
	HEMBA1003926	F-HEMBA1003926	564	R-HEMBA1003926	6090
	HEMBA1003937	F-HEMBA1003937	565	R-HEMBA1003937	6091
55	HEMBA1003939	F-HEMBA1003939	566	R-HEMBA1003939	6092
	HEMBA1003942	F-HEMBA1003942	567	R-HEMBA1003942	6093
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	HEMBA1003950	F-HEMBA1003950	568	R-HEMBA1003950	6094
	HEMBA1003953	F-HEMBA1003953	569	R-HEMBA1003953	6095
	HEMBA1003958	F-HEMBA1003958	570	R-HEMBA1003958	6096
5	HEMBA1003959	F-HEMBA1003959	571	R-HEMBA1003959	6097
	HEMBA1003976	F-HEMBA1003976	572	R-HEMBA1003976	6098
	HEMBA1003978	F-HEMBA1003978	573	R-HEMBA1003978	6099
	HEMBA1003985	F-HEMBA1003985	574	R-HEMBA1003985	6100
10	HEMBA1003987	F-HEMBA1003987	575	R-HEMBA1003987	6101
	HEMBA1003989	F-HEMBA1003989	576	R-HEMBA1003989	6102
	HEMBA1004000	F-HEMBA1004000	577	R-HEMBA1004000	6103
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15	HEMBA1004012	F-HEMBA1004012	579	R-HEMBA1004012	6105
	HEMBA1004015	F-HEMBA1004015	580	R-HEMBA1004015	6106
	HEMBA1004024	F-HEMBA1004024,	581	R-HEMBA1004024	6107
	HEMBA1004038	F-HEMBA1004038	582	R-HEMBA1004038	6108
20	HEMBA1004042	F-HEMBA1004042	583	R-HEMBA1004042	6109
20	HEMBA1004045	F-HEMBA1004045	584	R-HEMBA1004045	6110
	HEMBA1004048	F-HEMBA1004048	585	R-HEMBA1004048	6111
	HEMBA1004049	F-HEMBA1004049	586	R-HEMBA1004049	6112
0.5	HEMBA1004055	F-HEMBA1004055	587	R-HEMBA1004055	6113
25	HEMBA1004056	F-HEMBA1004056	588	R-HEMBA1004056	6114
	HEMBA1004074	F-HEMBA1004074	589	R-HEMBA1004074	6115
	HEMBA1004086	F-HEMBA1004086	590	R-HEMBA1004086	6116
	HEMBA1004097	F-HEMBA1004097	591	R-HEMBA1004097	6117
30	HEMBA1004111	F-HEMBA1004111	592		
	HEMBA1004131	F-HEMBA1004131	593	R-HEMBA1004131	6118
	HEMBA1004132	F-HEMBA1004132	594	R-HEMBA1004132	6119
	HEMBA1004133	F-HEMBA1004133	595	R-HEMBA1004133	6120
35	HEMBA1004138	F-HEMBA1004138	596	R-HEMBA1004138	6121
	HEMBA1004143	F-HEMBA1004143	597	R-HEMBA1004143	6122
	HEMBA1004146	F-HEMBA1004146	598	R-HEMBA1004146	6123
	HEMBA1004150	F-HEMBA1004150	599	R-HEMBA1004150	6124
40	HEMBA1004164	F-HEMBA1004164	600	R-HEMBA1004164	6125
	HEMBA1004168	F-HEMBA1004168	601	R-HEMBA1004168	6126
	HEMBA1004199	F-HEMBA1004199	602	R-HEMBA1004199	6127
	HEMBA1004200	F-HEMBA1004200	603	R-HEMBA1004200	6128
45	HEMBA1004202	F-HEMBA1004202	604	R-HEMBA1004202	6129
	HEMBA1004203	F-HEMBA1004203	605	R-HEMBA1004203	6130
	HEMBA1004207	F-HEMBA1004207	606	R-HEMBA1004207	6131
	HEMBA1004225	F-HEMBA1004225	607	R-HEMBA1004225	6132
50	HEMBA1004227	F-HEMBA1004227	608	R-HEMBA1004227	6133
	HEMBA1004238	F-HEMBA1004238	609	R-HEMBA1004238	6134
	HEMBA1004241	F-HEMBA1004241	610	R-HEMBA1004241	6135
	HEMBA1004246	F-HEMBA1004246	611	R-HEMBA1004246	6136
55	HEMBA1004248	F-HEMBA1004248	612	R-HEMBA1004248	6137
	HEMBA1004264	F-HEMBA1004264	613	R-HEMBA1004264	6138

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	HEMBA1004267	F-HEMBA1004267	614 ·	R-HEMBA1004267	6139
	HEMBA1004272	F-HEMBA1004272	615	R-HEMBA1004272	6140
	HEMBA1004274	F-HEMBA1004274	616		
5	HEMBA1004275	F-HEMBA1004275	617	R-nnnnnnnnnnn	6141
	HEMBA1004276	F-HEMBA1004276	618	R-HEMBA1004276	6142
	HEMBA1004286	F-HEMBA1004286	619	R-HEMBA1004286	6143
	HEMBA1004289	F-HEMBA1004289	620	R-HEMBA1004289	6144
10	HEMBA1004295	F-HEMBA1004295	621	R-HEMBA1004295	6145
	HEMBA1004306	F-HEMBA1004306	622	R-HEMBA1004306	6146
	HEMBA1004312	F-HEMBA1004312	623	R-HEMBA1004312	6147
	HEMBA1004321	F-HEMBA1004321	624	R-HEMBA1004321	6148
15	HEMBA1004323	F-HEMBA1004323	625	R-HEMBA1004323	6149
10	HEMBA1004327	F-HEMBA1004327	626	R-HEMBA1004327	6150
	HEMBA1004330	F-HEMBA1004330,	627	R-HEMBA1004330	6151
	HEMBA1004334	F-HEMBA1004334	628	R-HEMBA1004334	6152
•	HEMBA1004335	F-HEMBA1004335	629	R-HEMBA1004335	6153
20	HEMBA1004341	F-HEMBA1004341	630	R-HEMBA1004341	6154
	HEMBA1004353	F-HEMBA1004353	631	R-HEMBA1004353	6155
	HEMBA1004354	F-HEMBA1004354	632	R-HEMBA1004354	6156
	HEMBA1004356	F-HEMBA1004356	633	R-HEMBA1004356	6157
25	HEMBA1004366	F-HEMBA1004366	634	R-HEMBA1004366	6158
	HEMBA1004372	F-HEMBA1004372	635	R-HEMBA1004372	6159
	HEMBA1004389	F-HEMBA1004389	636	R-HEMBA1004389	6160
	HEMBA1004394	F-HEMBA1004394	637	R-HEMBA1004394	6161
30	HEMBA1004396	F-HEMBA1004396	638	R-HEMBA1004396	6162
	HEMBA1004405	F-HEMBA1004405	639	R-HEMBA1004405	6163
	HEMBA1004408	F-HEMBA1004408	640	R-HEMBA1004408	6164
	HEMBA1004429	F-HEMBA1004429	641	R-HEMBA1004429	6165
35	HEMBA1004433	F-HEMBA1004433	642	R-HEMBA1004433	6166
	HEMBA1004460	F-HEMBA1004460	643	R-HEMBA1004460	6167
	HEMBA1004461	F-HEMBA1004461	644	R-HEMBA1004461	6168
	HEMBA1004479	F-HEMBA1004479	645	R-HEMBA1004479	6169
40	HEMBA1004482	F-HEMBA1004482	646	R-HEMBA1004482	6170
	HEMBA1004499	F-HEMBA1004499	647	•	
	HEMBA1004502	F-HEMBA1004502	648	R-HEMBA1004502	6171
	HEMBA1004506	F-HEMBA1004506	649	R-HEMBA1004506	6172
45	HEMBA1004507	F-HEMBA1004507	650	R-HEMBA1004507	6173
	HEMBA1004509	F-HEMBA1004509	651	R-HEMBA1004509	6174
	HEMBA1004534	F-HEMBA1004534	652	R-HEMBA1004534	6175
	HEMBA1004538	F-HEMBA1004538	653	R-HEMBA1004538	6176
50	HEMBA1004542	F-HEMBA1004542	654		
	HEMBA1004554	F-HEMBA1004554	655	R-HEMBA1004554	6177
	HEMBA1004560	F-HEMBA1004560	656	R-HEMBA1004560	6178
	HEMBA1004573	F-HEMBA1004573	657	R-HEMBA1004573	6179
55	HEMBA1004577	F-HEMBA1004577	658	R-HEMBA1004577	6180
	HEMBA1004586	F-HEMBA1004586	659	R-HEMBA1004586	6181

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	HEMBA1004596	F-HEMBA1004596	660	R-nnnnnnnnnnn	6182
	HEMBA1004604	F-HEMBA1004604	661		
	HEMBA1004610	F-HEMBA1004610	662	R-HEMBA1004610	6183
5	HEMBA1004617	F-HEMBA1004617	663	R-HEMBA1004617	6184
	HEMBA1004629	F-HEMBA1004629	664	R-HEMBA1004629	6185
	HEMBA1004631	F-HEMBA1004631	665	R-HEMBA1004631	6186
	HEMBA1004632	F-HEMBA1004632	666	R-HEMBA1004632	6187
10	HEMBA1004637	F-HEMBA1004637	667	R-HEMBA1004637	6188
	HEMBA1004638	F-HEMBA1004638	668	R-HEMBA1004638	6189
	HEMBA1004666	F-HEMBA1004666	669	R-HEMBA1004666	6190
	HEMBA1004669	F-HEMBA1004669	670	R-HEMBA1004669	6191
15	HEMBA1004670	F-HEMBA1004670	671	R-HEMBA1004670	6192
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	HEMBA1004697	F-HEMBA1004697	674	R-HEMBA1004697	6195
20	HEMBA1004705	F-HEMBA1004705	675	R-HEMBA1004705	6196
20	HEMBA1004709	F-HEMBA1004709	676	R-HEMBA1004709	6197
	HEMBA1004711	F-HEMBA1004711	677	R-HEMBA1004711	6198
	HEMBA1004725	F-HEMBA1004725	678	R-HEMBA1004725	6199
	HEMBA1004730	F-HEMBA1004730	679	R-HEMBA1004730	6200
25	HEMBA1004733	F-HEMBA1004733	680	R-HEMBA1004733	6201
	HEMBA1004734	F-HEMBA1004734	681	R-HEMBA1004734	6202
	HEMBA1004736	F-HEMBA1004736	682	R-HEMBA1004736	6203
	HEMBA1004748	F-HEMBA1004748	683	R-HEMBA1004748	6204
30	HEMBA1004751	F-HEMBA1004751	684	R-HEMBA1004751	6205
	HEMBA1004752	F-HEMBA1004752	685	R-HEMBA1004752	6206
	HEMBA1004753	F-HEMBA1004753	686	R-HEMBA1004753	6207
	HEMBA1004756	F-HEMBA1004756	687	R-HEMBA1004756	6208
35	HEMBA1004758	F-HEMBA1004758	688	R-HEMBA1004758	6209
	HEMBA1004763	F-HEMBA1004763	689	R-HEMBA1004763	6210
	HEMBA1004768	F-HEMBA1004768	690	R-HEMBA1004768	6211
	HEMBA1004770	F-HEMBA1004770	691	R-HEMBA1004770	6212
40	HEMBA1004771	F-HEMBA1004771	692	R-HEMBA1004771	6213
	HEMBA1004776	F-HEMBA1004776	693	R-HEMBA1004776	6214
	HEMBA1004778	F-HEMBA1004778	694	R-HEMBA1004778	6215
	HEMBA1004795	F-HEMBA1004795	695	R-nananananan	6216
45	HEMBA1004803	F-HEMBA1004803	696	R-HEMBA1004803	6217
,,,	HEMBA1004806	F-HEMBA1004806	697	R-HEMBA1004806	6218
	HEMBA1004807	F-HEMBA1004807	698	R-HEMBA1004807	6219
	HEMBA1004816	F-HEMBA1004816	699	R-HEMBA1004816	6220
50	HEMBA1004820	F-HEMBA1004820	700	R-HEMBA1004820	6221
30	HEMBA1004847	F-HEMBA1004847	701	R-HEMBA1004847	6222
	HEMBA1004850	F-HEMBA1004850	702	R-HEMBA1004850	6223
	HEMBA1004863	F-HEMBA1004863	703	R-HEMBA1004863	6224
	HEMBA1004864	F-HEMBA1004864	704	R-HEMBA1004864	6225
55	HEMBA1004865	F-HEMBA1004865	705	R-HEMBA1004865	6226

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	706 R-HEMBA100400
	HEMBA1004880 F-HEMBA1004889 6228
	HEMBA1004889 F-HEMBA1004900 6229
	HEMBA1004900 F-HEMBA1004909 0230
	HEMBA1004909 F-REMBA1004918 0231
5	HEMBA1004918 F-HEMBA1004923 6232
	HEMBA1004923 F-HEMBA1004929 6233
	HEMBA1004929 F-HEMBA1004930 0234
	HEMBA1004930 F-HEMBA1004933 6235
10	HEMBA1004933 F-HEMBA1004934 0230
	HEMBA1004934 F-HEMBA1004934 716 R-HEMBA1004944 6237
	HEMBA1004944 F-HEMBA1004954 0230
	HEMBA1004954 F-HEMBA1004954 718 R-HEMBA1004956 6239
15	HEMBA1004956 F-HEMBA1004960 0240
	HEMBA1004960 F-HEMBA1004900 720 R-HEMBA1004972 6241
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	1005050 F-HEMBA1005050 734 R-1111105062 6256
35	WEIGHT 1005062 F-HEMBA1005062 735 RELEASE 1005066 6257
	1005066 F-HEMBA1000000 - 5 10000105075 6250
	TEMPA 1 005075 F-HEMBA 1005070
	1005079 F-HEMBA1005015 1005083 6260
40	HEMBA1005083 F-HEMBA1005003 7/0 R-HEMBA1005101 6201
	HEMBA1005101 F-HEMBA1005101 741 R-HEMBA1005113 6262
	HEMBA1005113 F-HEMBA1005113 742 R-HEMBA1005123 6263
	HEMBA1005123 F-HEMBA1005123 R-HEMBA1005133 6264
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	HEMBA1005152 F-HEMBA1005159 0201
	HEMBA1005159 F-HEMBA1005185 0200
50	HEMBA1005185 F-HEMBA1005201 0205
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55	HEMBA1005219 F-HEMBA1003213

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	HEMBA1005223	F-HEMBA1005223	752	R-HEMBA1005223	6272
	HEMBA1005232	F-HEMBA1005232	753	R-HEMBA1005232	6273
	HEMBA1005241	F-HEMBA1005241	754	R-HEMBA1005241	6274
5	HEMBA1005244	F-HEMBA1005244	755	R-HEMBA1005244	6275
	HEMBA1005251	F-HEMBA1005251	756	R-HEMBA1005251	6276
	HEMBA1005252	F-HEMBA1005252	757	R-HEMBA1005252	6277
	HEMBA1005274	F-HEMBA1005274	758	R-HEMBA1005274	6278
10	HEMBA1005275	F-HEMBA1005275	759	R-HEMBA1005275	6279
10	HEMBA1005293	F-HEMBA1005293	760	R-HEMBA1005293	6280
	HEMBA1005296	F-HEMBA1005296	761	R-HEMBA1005296	6281
	HEMBA1005304	F-HEMBA1005304	762	R-HEMBA1005304	6282
	HEMBA1005311	F-HEMBA1005311	763	R-HEMBA1005311	6283
15	HEMBA1005314	F-HEMBA1005314	764	R-HEMBA1005314	6284
	HEMBA1005315	F-HEMBA1005315,	765	R-HEMBA1005315	6285
	HEMBA1005318	F-HEMBA1005318	766	R-HEMBA1005318	6286
	HEMBA1005331	F-HEMBA1005331	767	R-HEMBA1005331	6287
20	HEMBA1005338	F-HEMBA1005338	768	N ILLEDATOUSSI	0201
	HEMBA1005353	F-HEMBA1005353	769	R-HEMBA1005353	6288
	HEMBA1005359	F-HEMBA1005359	770	R-HEMBA1005359	6289
	HEMBA1005367	F-HEMBA1005367	771	R-HEMBA1005367	6290
25	HEMBA1005372	F-HEMBA1005372	772	R-HEMBA1005372	6291
	HEMBA1005374	F-HEMBA1005374	773	R-HEMBA1005374	6292
	HEMBA1005382	F-HEMBA1005382	774	W HEMOWICON	0232
	HEMBA1005389	F-HEMBA1005389	775	R-HEMBA1005389	6293
30	HEMBA1005394	F-HEMBA1005394	776	R-HEMBA1005394	6294
50	HEMBA1005403	F-HEMBA1005403	777	R-HEMBA1005403	6295
	HEMBA1005408	F-HEMBA1005408	778	R-HEMBA1005408	6296
	HEMBA1005400	F-HEMBA1005410	779	R-HEMBA1005410	6297
25	HEMBA1005411	F-HEMBA1005411	780	R-HEMBA1005411	6298
35	HEMBA1005411	F-HEMBA1005423	781	R-HEMBA1005423	6299
	HEMBA1005425	F-HEMBA1005426	782	R-HEMBA1005426	6300
	HEMBA1005443	F-HEMBA1005443	783	R-HEMBA1005443	6301
	HEMBA1005447	F-HEMBA1005447	784	R-HEMBA1005447	6302
40	HEMBA1005468	F-HEMBA1005468	785	R-HEMBA1005468	6303
	HEMBA1005469	F-HEMBA1005469	786		6304
	HEMBA1005472	F-HEMBA1005472	787	R-HEMBA1005472	6305
	HEMBA1005474	F-HEMBA1005474	788	R THEMBATOUS 412	0303
45	HEMBA1005475	F-HEMBA1005475	789	R-HEMBA1005475	6306
	HEMBA1005497		790	R-HEMBA1005497	6307
	HEMBA1005500	F-HEMBA1005500	791	R-HEMBA1005500	6308
	HEMBA1005506	F-HEMBA1005506	792	R-HEMBA1005506	6309
50	HEMBA1005508	F-HEMBA1005508	793	R-HEMBA1005508	6310
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	HEMBA1005511	F-HEMBA1005511 F-HEMBA1005513	794 795	V_UEWDV100911	6311
	HEMBA1005513			R-HEMBA1005517	6210
55	HEMBA1005517	F-HEMBA1005517	796 707		6312
-	HEMBA1005518	F-HEMBA1005518	797	R-HEMBA1005518	6313

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	HEMBA1005520	F-HEMBA1005520	798	R-HEMBA1005520	6314
	HEMBA1005526	F-HEMBA1005526	799	R-HEMBA1005526	6315
	HEMBA1005528	F-HEMBA1005528	800	R-HEMBA1005528	6316
5	HEMBA1005530	F-HEMBA1005530	801	R-HEMBA1005530	6317
	HEMBA1005548	F-HEMBA1005548	802	R-HEMBA1005548	6318
	HEMBA1005552	F-HEMBA1005552	803	R-HEMBA1005552	6319
	HEMBA1005558	F-HEMBA1005558	804	R-HEMBA1005558	6320
10	HEMBA1005568	F-HEMBA1005568	805	R-HEMBA1005568	6321
,•	HEMBA1005570	F-HEMBA1005570	806	R-HEMBA1005570	6322
	HEMBA1005576	F-HEMBA1005576	807	R-HEMBA1005576	6323
	HEMBA1005577	F-HEMBA1005577	808	R-HEMBA1005577	6324
15	HEMBA1005581	F-HEMBA1005581	809	R-HEMBA1005581	6325
15	HEMBA1005582	F-HEMBA1005582	810	R-HEMBA1005582	6326
	HEMBA1005583	F-HEMBA1005583,	811	R-HEMBA1005583	6327
	HEMBA1005588	F-HEMBA1005588	812	R-HEMBA1005588	6328
	HEMBA1005593	F-HEMBA1005593	813	R-HEMBA1005593	6329
20	HEMBA1005595	F-HEMBA1005595	814	R-HEMBA1005595	6330
	HEMBA1005606	F-HEMBA1005606	815	R-HEMBA1005606	6331
	HEMBA1005609	F-HEMBA1005609	816	R-HEMBA1005609	6332
	HEMBA1005616	F-HEMBA1005616	817	R-HEMBA1005616	6333
25	HEMBA1005621	F-HEMBA1005621	818	R-HEMBA1005621	6334
	HEMBA1005627	F-HEMBA1005627	819	R-HEMBA1005627	6335
	HEMBA1005631	F-HEMBA1005631	820	R-HEMBA1005631	6336
	HEMBA1005632	F-HEMBA1005632	821	R-HEMBA1005632	6337
30	HEMBA1005634	F-HEMBA1005634	822	R-HEMBA1005634	6338
	HEMBA1005666	F-HEMBA1005666	823	R-HEMBA1005666	6339
	HEMBA1005670	F-HEMBA1005670	824	R-HEMBA1005670	6340
	HEMBA1005679	F-HEMBA1005679	825	R-HEMBA1005679	6341
35	HEMBA1005680	F-HEMBA1005680	826	R-HEMBA1005680	6342
	HEMBA1005685	F-HEMBA1005685	827	R-HEMBA1005685	6343
	HEMBA1005699	F-HEMBA1005699	828	R-HEMBA1005699	6344
	HEMBA1005705	F-HEMBA1005705	829	R-HEMBA1005705	6345
40	HEMBA1005717	F-HEMBA1005717	830	R-HEMBA1005717	6346
	HEMBA1005732	F-HEMBA1005732	831	R-HEMBA1005732	6347
	HEMBA1005737	F-HEMBA1005737	832	R-HEMBA1005737	6348
	HEMBA1005746	F-HEMBA1005746	833	R-nnnnnnnnnnn	6349
45	HEMBA1005755	F-HEMBA1005755	834	R-HEMBA1005755	6350
	HEMBA1005765	F-HEMBA1005765	835	R-HEMBA1005765	6351
	HEMBA1005780	F-HEMBA1005780	836	R-HEMBA1005780	6352
	HEMBA1005813	F-HEMBA1005813	837	R-HEMBA1005813	6353
50	HEMBA1005815	F-HEMBA1005815	838	R-HEMBA1005815	6354
30	HEMBA1005822	F-HEMBA1005822	839	R-HEMBA1005822	6355
	HEMBA1005829	F-HEMBA1005829	840	R-HEMBA1005829	6356
	HEMBA1005834	F-HEMBA1005834	841	R-HEMBA1005834	6357
F.C.	HEMBA1005852	F-HEMBA1005852	842	R-HEMBA1005852	6358
55	HEMBA1005853	F-HEMBA1005853	843	R-HEMBA1005853	6359

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	HEMBA1005884	F-HEMBA1005884	844	R-HEMBA1005884	6360
	HEMBA1005891	F-HEMBA1005891	845	R-HEMBA1005891	6361
	HEMBA1005894	F-HEMBA1005894	846	R-HEMBA1005894	6362
5	HEMBA1005909	F-HEMBA1005909	847	R-HEMBA1005909	6363
	HEMBA1005911	F-HEMBA1005911	848	R-HEMBA1005911	6364
	HEMBA1005921	F-HEMBA1005921	849	R-HEMBA1005921	6365
	HEMBA1005931	F-HEMBA1005931	850	R-HEMBA1005931	6366
10	HEMBA1005934	F-HEMBA1005934	851	R-HEMBA1005934	6367
10	HEMBA1005962	F-HEMBA1005962	852	R-HEMBA1005962	6368
	HEMBA1005963	F-HEMBA1005963	853	R-HEMBA1005963	6369
	HEMBA1005990	F-HEMBA1005990	854	R-HEMBA1005990	6370
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15	HEMBA1005999	F-HEMBA1005999	856	R-HEMBA1005999	6372
	HEMBA1006002	F-HEMBA1006002,	857	R-HEMBA1006002	6373
	HEMBA1006005	F-HEMBA1006005	858	R-HEMBA1006005	6374
	HEMBA1006031	F-HEMBA1006031	859	R-nnnnnnnnnnn	6375
20	HEMBA1006035	F-HEMBA1006035	860	R-HEMBA1006035	6376
	HEMBA1006036	F-HEMBA1006036	861	R-HEMBA1006036	6377
	HEMBA1006042	F-HEMBA1006042	862	R-HEMBA1006042	6378
	HEMBA1006067	F-HEMBA1006067	863	R-nnnnnnnnnnn	6379
25	HEMBA1006081	F-HEMBA1006081	864	R-HEMBA1006081	6380
	HEMBA1006090	F-HEMBA1006090	865	R-HEMBA1006090	6381
	HEMBA1006090	F-HEMBA1006090	866	R-HEMBA1006091	6382
	HEMBA1006100	F-HEMBA1006100	867	R-HEMBA1006100	6383
30	HEMBA1006108	F-HEMBA1006108	868	R-HEMBA1006108	6384
	HEMBA1006121	F-HEMBA1006121	869	R-HEMBA1006121	6385
,	HEMBA1006124	F-HEMBA1006124	870	R-HEMBA1006124	6386
	HEMBA1006130	F-HEMBA1006130	871	R-HEMBA1006130	6387
25	HEMBA1006138	F-HEMBA1006138	872	R-nnnnnnnnnnn	6388
35	HEMBA1006142	F-HEMBA1006142	873	R-HEMBA1006142	6389
	HEMBA1006155	F-HEMBA1006155	874	R-HEMBA1006155	6390
	HEMBA1006158	F-HEMBA1006158	875	R-HEMBA1006158	6391
	HEMBA1006173	F-HEMBA1006173	876	R-HEMBA1006173	6392
40	HEMBA1006182	F-HEMBA1006182	877	R-HEMBA1006182	6393
	HEMBA1006198	F-HEMBA1006198	878	R-HEMBA1006198	6394
	HEMBA1006235	F-HEMBA1006235	879	R-HEMBA1006235	6395
	HEMBA1006248	F-HEMBA1006248	880	R-HEMBA1006248	6396
45	HEMBA1006252	F-HEMBA1006252	881	R-HEMBA1006252	6397
	HEMBA1006253	F-HEMBA1006253	882	R-HEMBA1006253	6398
	HEMBA1006259	F-HEMBA1006259	883	R-HEMBA1006259	6399
	HEMBA1006268	F-HEMBA1006268	884	R-HEMBA1006268	6400
50	HEMBA1006272	F-HEMBA1006272	885	R-HEMBA1006272	6401
		F-HEMBA1006278	886	R-nnnnnnnnnnn	6402
	HEMBA1006278	F-HEMBA1006283	887	R-HEMBA1006283	6403
	HEMBA1006283		888	R-HEMBA1006284	6404
55	HEMBA1006284	F-HEMBA1006284 F-HEMBA1006291	889	R-HEMBA1006291	6405
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	HEMBA1006309	F-HEMBA1006309	891	R-HEMBA1006309	6407
	HEMBA1006310	F-HEMBA1006310	892	R-HEMBA1006310	6408
5	HEMBA1006328	F-HEMBA1006328	893	R-HEMBA1006328	6409
	HEMBA1006334	F-HEMBA1006334	894	R-HEMBA1006334	6410
	HEMBA1006344	F-HEMBA1006344	895	R-HEMBA1006344	6411
	HEMBA1006347	F-HEMBA1006347	896	R-HEMBA1006347	6412
10	HEMBA1006349	F-HEMBA1006349	897	R-HEMBA1006349	6413
70	HEMBA1006359	F-HEMBA1006359	898	R-HEMBA1006359	6414
	HEMBA1006364	F-HEMBA1006364	899	R-HEMBA1006364	6415
	HEMBA1006377	F-HEMBA1006377	900	R-HEMBA1006377	6416
	HEMBA1006380	F-HEMBA1006380	901	R-HEMBA1006380	6417
15	HEMBA1006381	F-HEMBA1006381	902	R-HEMBA1006381	6418
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20	HEMBA1006421	F-HEMBA1006421	906	R-HEMBA1006421	6422
	HEMBA1006424	F-HEMBA1006424	907	R-HEMBA1006424	6423
	HEMBA1006426	F-HEMBA1006426	908	R-HEMBA1006426	6424
	HEMBA1006438	F-HEMBA1006438	909	R-HEMBA1006438	6425
25	HEMBA1006445	F-HEMBA1006445	910	R-HEMBA1006445	6426
	HEMBA1006446	F-HEMBA1006446	911	R-HEMBA1006446	6427
	HEMBA1006461	F-HEMBA1006461	912	R-HEMBA1006461	6428
	HEMBA1006467	F-HEMBA1006467	913	R-HEMBA1006467	6429
30	HEMBA1006471	F-HEMBA1006471	914	R-HEMBA1006471	6430
	HEMBA1006474	F-HEMBA1006474	915	R-HEMBA1006474	6431
,	HEMBA1006483	F-HEMBA1006483	916	R-HEMBA1006483	6432
	HEMBA1006485	F-HEMBA1006485	917	R-HEMBA1006485	6433
35	HEMBA1006486	F-HEMBA1006486	918	R-HEMBA1006486	6434
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	HEMBA1006492	F-HEMBA1006492	920	R-HEMBA1006492	6436
	HEMBA1006494	F-HEMBA1006494	921	R-HEMBA1006494	6437
40	HEMBA1006497	F-HEMBA1006497	922	R-HEMBA1006497	6438
40	HEMBA1006502	F-HEMBA1006502	923	R-HEMBA1006502	6439
	HEMBA1006507	F-HEMBA1006507	924	R-HEMBA1006507	6440
	HEMBA1006521	F-HEMBA1006521	925	R-HEMBA1006521	6441
	HEMBA1006530	F-HEMBA1006530	926	R-HEMBA1006530	6442
45	HEMBA1006535	F-HEMBA1006535	927	R-HEMBA1006535	6443
	HEMBA1006540	F-HEMBA1006540	928	R-HEMBA1006540	6 44 4
	HEMBA1006546	F-HEMBA1006546	929	R-HEMBA1006546	6445
	HEMBA1006559	F-HEMBA1006559	930	R-HEMBA1006559	6446
50	HEMBA1006562	F-HEMBA1006562	931	R-HEMBA1006562	6447
	HEMBA1006566	F-HEMBA1006566	932	R-HEMBA1006566	6448
	HEMBA1006569	F-HEMBA1006569	933	R-HEMBA1006569	6449
	HEMBA1006579	F-HEMBA1006579	934	R-HEMBA1006579	6450
55	HEMBA1006583	F-HEMBA1006583	935	R-HEMBA1006583	6451
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	HEMBA1006595	F-HEMBA1006595	936	R-HEMBA1006595	6452
	HEMBA1006597	F-HEMBA1006597	937	R-HEMBA1006597	6453
	HEMBA1006612	F-HEMBA1006612	938	R-HEMBA1006612	6454
5	HEMBA1006617	F-HEMBA1006617	939	R-nnnnnnnnnn	6455
	HEMBA1006624	F-HEMBA1006624	940	R-HEMBA1006624	6456
	HEMBA1006631	F-HEMBA1006631	941	R-HEMBA1006631	6457
	HEMBA1006635	F-HEMBA1006635	942	R-HEMBA1006635	6458
	HEMBA1006639	F-HEMBA1006639	943	R-HEMBA1006639	6459
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	HEMBA1006652	F-HEMBA1006652	946	R-HEMBA1006652	6462
	HEMBA1006653	F-HEMBA1006653	947	R-HEMBA1006653	6463
15	HEMBA1006659		948	K IEMPYTOOOOO	0400
		F-HEMBA1006659		D DEMONTORESE	6464
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	HEMBA1006674	F-HEMBA1006674	950	R-HEMBA1006674	
20	HEMBA1006676	F-HEMBA1006676	951	R-HEMBA1006676	6466
	HEMBA1006682	F-HEMBA1006682	952	R-HEMBA1006682	6467
	HEMBA1006695	F-HEMBA1006695	953	R-HEMBA1006695	6468
	HEMBA1006696	F-HEMBA1006696	954	R-HEMBA1006696	6469
25	HEMBA1006708	F-HEMBA1006708	955	R-HEMBA1006708	6470
20	HEMBA1006709	F-HEMBA1006709	956	R-HEMBA1006709	6471
	HEMBA1006717	F-HEMBA1006717	957	R-HEMBA1006717	6472
	HEMBA1006737	F-HEMBA1006737	958	R-HEMBA1006737	6473
	HEMBA1006744	F-HEMBA1006744	959	R-HEMBA1006744	6474
30	HEMBA1006754	F-HEMBA1006754	960	R-HEMBA1006754	6475
	HEMBA1006758	F-HEMBA1006758	961	R-HEMBA1006758	6476
	HEMBA1006767	F-HEMBA1006767	962	R-HEMBA1006767	6477
	HEMBA1006779	F-HEMBA1006779	963	R-HEMBA1006779	6478
35	HEMBA1006780	F-HEMBA1006780	964	R-HEMBA1006780	6479
	HEMBA1006789	F-HEMBA1006789	965	R-HEMBA1006789	6480
	HEMBA1006795	F-HEMBA1006795	966	R-HEMBA1006795	6481
	HEMBA1006796	F-HEMBA1006796	967	R-HEMBA1006796	6482
40	HEMBA1006807	F-HEMBA1006807	968	R-HEMBA1006807	6483
	HEMBA1006821	F-HEMBA1006821	969	R-HEMBA1006821	6484
	HEMBA1006824	F-HEMBA1006824	970	R-HEMBA1006824	6485
	HEMBA1006832	F-HEMBA1006832	971	R-HEMBA1006832	6486
45	HEMBA1006849	F-HEMBA1006849	972	R-HEMBA1006849	6487
	HEMBA1006865	F-HEMBA1006865	973	R-HEMBA1006865	6488
	HEMBA1006877	F-HEMBA1006877	974	R-nnnnnnnnnnn	6489
	HEMBA1006885	F-HEMBA1006885	975	R-HEMBA1006885	6490
50	HEMBA1006900	F-HEMBA1006900	976	R-HEMBA1006900	6491
50	HEMBA1006914	F-HEMBA1006914	977		
	HEMBA1006921	F-HEMBA1006921	978	R-HEMBA1006921	6492
	HEMBA1006926	F-HEMBA1006926	979	R-HEMBA1006926	6493
	HEMBA1006929	F-HEMBA1006929	980	R-HEMBA1006929	6494
55	HEMBA1006936	F-HEMBA1006936	981	R-HEMBA1006936	6495
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	HEMBA1006938	F-HEMBA1006938	982	R-HEMBA1006938	6496
	HEMBA1006941	F-HEMBA1006941	983	R-HEMBA1006941	6497
	HEMBA1006949	F-HEMBA1006949	984	R-HEMBA1006949	6498
5	HEMBA1006973	F-HEMBA1006973	985	R-HEMBA1006973	6499
	HEMBA1006976	F-HEMBA1006976	986	R-HEMBA1006976	6500
	HEMBA1006993	F-HEMBA1006993	987	R-HEMBA1006993	6501
	HEMBA1006996	F-HEMBA1006996	988	R-HEMBA1006996	6502
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10	HEMBA1007017	F-HEMBA1007017	990	R-HEMBA1007017	6504
	HEMBA1007018	F-HEMBA1007018	991	R-HEMBA1007018	6505
	HEMBA1007045	F-HEMBA1007045	992	R-HEMBA1007045	6506
	HEMBA1007051	F-HEMBA1007051	993	R-HEMBA1007051	6507
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	HEMBA1007052	F-HEMBA1007062,	995	R-HEMBA1007062	6509
	HEMBA1007066	F-HEMBA1007066	996	R-HEMBA1007066	6510
	HEMBA1007073	F-HEMBA1007073	997	R-HEMBA1007073	6511
20	НЕМВА1007073 НЕМВА1007078	F-HEMBA1007078	998	R-HEMBA1007078	6512
	немви 1007078 НЕМВА 1007080	F-HEMBA1007080	999	K-UEMDY1001019	0312
		F-HEMBA1007085	1000	R-HEMBA1007085	6513
	HEMBA1007085 HEMBA1007087	F-HEMBA1007087	1000	R-HEMBA1007087	6514
25	HEMBA1007087	F-HEMBA1007037	1001	R-HEMBA1007112	6515
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		F-HEMBA1007121	1003	K-HEMDATOV1113	0010
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•	HEMBA1007147	F-HEMBA1007147	1005	R-HEMBA1007147	6518
30	HEMBA1007147	F-HEMBA1007147	1007	R-HEMBA1007149	6519
	HEMBA1007151	F-HEMBA1007151	1007	R-HEMBA1007151	6520
	HEMBA1007174	F-HEMBA1007174	1008	R-nnnnnnnnnnn	6521
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35	HEMBA1007178	F-HEMBA1007194	1010	R-HEMBA1007194	6523
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	HEMBA1007206	F-HEMBA1007206	1013	R-HEMBA1007206	6525
	HEMBA1007224	F-HEMBA1007224	1013	R-HEMBA1007224	6526
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	HEMBA1007251	F-HEMBA1007251	1016	R-HEMBA1007251	6527
	HEMBA1007256	F-HEMBA1007256	1017	R-HEMBA1007256	6528
	HEMBA1007267	F-HEMBA1007267	1018	R-HEMBA1007267	6529
45	HEMBA1007273	F-HEMBA1007273	1019	R-HEMBA1007273	6530
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30	HEMBA1007301	F-HEMBA1007301	1024	R-HEMBA1007301	6535
	HEMBA1007319	F-HEMBA1007319	1025	R-HEMBA1007319	6536
	HEMBA1007319	F-HEMBA1007320	1026	R-HEMBA1007320	6537
	HEMBA1007322	F-HEMBA1007322	1027	R-HEMBA1007322	6538
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	HEMBA1007341	F-HEMBA1007341	1029	R-HEMBA1007341	6540
	HEMBA1007342	F-HEMBA1007342	1030	R-HEMBA1007342	6541
5	HEMBA1007347	F-HEMBA1007347	1031	R-HEMBA1007347	6542
-	HEMBB1000005	F-HEMBB1000005	1032	R-HEMBB1000005	6543
	HEMBB1000008	F-HEMBB1000008	1033	R-HEMBB1000008	6544
	HEMBB1000018	F-HEMBB1000018	1034	R-HEMBB1000018	6545
10	HEMBB1000024	F-HEMBB1000024	1035	R-HEMBB1000024	6546
	HEMBB1000025	F-HEMBB1000025	1036	R-HEMBB1000025	6547
	HEMBB1000030	F-HEMBB1000030	1037	R-HEMBB1000030	6548
	HEMBB1000036	F-HEMBB1000036	1038	R-HEMBB1000036	6549
	HEMBB1000037	F-HEMBB1000037	1039	R-HEMBB1000037	6550
15	HEMBB1000039	F-HEMBB1000039	1040	R-HEMBB1000039	6551
	HEMBB1000044	F-HEMBB1000044,	1041	R-HEMBB1000044	6552
	HEMBB1000048	F-HEMBB1000048	1042	R-HEMBB1000048	6553
	HEMBB1000050	F-HEMBB1000050	1043	R-HEMBB1000050	6554
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	HEMBB1000055	F-HEMBB1000055	1045	R-HEMBB1000055	6556
	HEMBB1000059	F-HEMBB1000059	1046	R-HEMBB1000059	6557
	HEMBB1000083	F-HEMBB1000083	1047	R-HEMBB1000083	6558
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	HEMBB1000099	F-HEMBB1000099	1049	R-HEMBB1000099	6560
	HEMBB1000103	F-HEMBB1000103	1050	R-HEMBB1000103	6561
	HEMBB1000113	F-HEMBB1000113	1051	R-HEMBB1000113	6562
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	HEMBB1000141	F-HEMBB1000141	1054	R-HEMBB1000141	6565
	HEMBB1000144	F-HEMBB1000144	1055	R-HEMBB1000144	6566
35	HEMBB1000173	F-HEMBB1000173	1056	R-HEMBB1000173	6567
	HEMBB1000175	F-HEMBB1000175	1057	R-HEMBB1000175	6568
	HEMBB1000198	F-HEMBB1000198	1058	R-HEMBB1000198	6569
	HEMBB1000215	F-HEMBB1000215	1059	R-HEMBB1000215	6570
40	HEMBB1000217	F-HEMBB1000217	1060	R-HEMBB1000217	6571
40	HEMBB1000218	F-HEMBB1000218	1061	R-HEMBB1000218	6572
	HEMBB1000226	F-HEMBB1000226	1062	R-HEMBB1000226	6573
	HEMBB1000240	F-HEMBB1000240	1063	R-HEMBB1000240	6574
	HEMBB1000244	F-HEMBB1000244	1064	R-HEMBB1000244	6575
45	HEMBB1000250	F-HEMBB1000250	1065	R-HEMBB1000250	6576
	HEMBB1000258	F-HEMBB1000258	1066	R-HEMBB1000258	6577
	HEMBB1000264	F-HEMBB1000264	1067	R-HEMBB1000264	6578
	HEMBB1000266	F-HEMBB1000266	1068	R-HEMBB1000266	6579
50	HEMBB1000272	F-HEMBB1000272	1069	R-HEMBB1000272	6580
	HEMBB1000274	F-HEMBB1000274	1070	R-HEMBB1000274	6581
	HEMBB1000284	F-HEMBB1000284	1071	R-HEMBB1000284	6582
	HEMBB1000307	F-HEMBB1000307	1072	R-HEMBB1000307	6583
55	HEMBB1000312	F-HEMBB1000312	1073	R-HEMBB1000312	6584

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	HEMBB1000317	F-HEMBB1000317	1074	R-HEMBB1000317	6585
	HEMBB1000318	F-HEMBB1000318	1075	R-HEMBB1000318	6586
	HEMBB1000335	F-HEMBB1000335	1076	R-HEMBB1000335	6587
5	HEMBB1000336	F-HEMBB1000336	1077	R-HEMBB1000336	6588
	HEMBB1000337	F-HEMBB1000337	1078	R-HEMBB1000337	6589
	HEMBB1000338	F-HEMBB1000338	1079	R-HEMBB1000338	6590
	HEMBB1000339	F-HEMBB1000339	1080	R-HEMBB1000339	6591
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70	HEMBB1000343	F-HEMBB1000343	1082	R-HEMB81000343	6593
	HEMBB1000354	F-HEMBB1000354	1083	R-HEMBB1000354	6594
	HEMBB1000369	F-HEMBB1000369	1084	R-HEMBB1000369	6595
	HEMBB1000374	F-HEMBB1000374	1085	R-HEMBB1000374	6596
15	HEMBB1000376	F-HEMBB1000376	1086	R-HEMBB1000376	6597
	HEMBB1000391	F-HEMBB1000391,	1087	R-HEMBB1000391	6598
	HEMBB1000399	F-HEMBB1000399	1088	R-HEMBB1000399	6599
	HEMBB1000402	F-HEMBB1000402	1089	R-HEMBB1000402	6600
20	HEMBB1000404	F-HEMBB1000404	1090	R-HEMBB1000402	6601
	HEMBB1000420	F-HEMBB1000420	1091	R-HEMBB1000420	6602
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	HEMBB1000438	F-HEMBB1000438	1093	R-HEMBB1000438	6604
25	HEMBB1000441	F-HEMBB1000441	1094	R-HEMBB1000441	6605
	HEMBB1000449	F-HEMBB1000449	1095	R-HEMBB1000449	6606
	HEMBB1000455	F-HEMBB1000455	1096	R-HEMBB1000455	6607
	HEMBB1000472	F-HEMBB1000472	1097	R-HEMBB1000472	6608
30	HEMBB1000480	F-HEMBB1000480	1098	R-HEMBB1000480	6609
	HEMBB1000487	F-HEMBB1000487	1099	R-HEMBB1000487	6610
	HEMBB1000490	F-HEMBB1000490	1100	R-HEMBB1000490	6611
	HEMBB1000491	F-HEMBB1000491	1101	R-HEMBB1000491	6612
25	HEMBB1000493	F-HEMBB1000493	1102	R-HEMBB1000493	6613
35	HEMBB1000510	F-HEMB81000510	1103	R-HEMBB1000510	6614
	HEMBB1000518	F-HEMBB1000518	1104	R-HEMBB1000518	6615
	HEMBB1000523	F-HEMBB1000523	1105	R-HEMBB1000523	6616
	HEMBB1000530	F-HEMBB1000530	1106	R-HEMBB1000530	6617
40	HEMBB1000550	F-HEMBB1000550	1107	R-HEMBB1000550	6618
	HEMBB1000554	F-HEMBB1000554	1108	R-HEMBB1000554	6619
	HEMBB1000556	F-HEMBB1000556	1109	R-HEMBB1000556	6620
	HEMBB1000564	F-HEMBB1000564	1110	R-HEMBB1000564	6621
45	HEMBB1000573	F-HEMBB1000573	1111	R-HEMBB1000573	6622
	HEMBB1000575	F-HEMBB1000575	1112	R-HEMBB1000575	6623
	HEMBB1000586	F-HEMBB1000586	1113	R-HEMBB1000586	6624
	HEMBB1000589	F-HEMBB1000589	1114	R-HEMBB1000589	6625
50	HEMBB1000591	F-HEMBB1000591	1115	R-HEMBB1000591	6626
	HEMBB1000592	F-HEMBB1000592	1116	R-HEMBB1000592	6627
	HEMBB1000593	F-HEMBB1000593	1117		
	HEMBB1000598	F-HEMBB1000598	1118	R-HEMBB1000598	6628
55	HEMBB1000623	F-HEMBB1000623	1119	R-HEMBB1000623	6629

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	HEMBB1000630	F-HEMBB1000630	1120	R-HEMBB1000630	6630	
	HEMBB1000631	F-HEMBB1000631	1121	R-HEMBB1000631	6631	
	HEMBB1000632	F-HEMBB1000632	1122	R-HEMBB1000632	6632	
5	HEMBB1000637	F-HEMBB1000637	1123	R-HEMBB1000637	6633	
	HEMBB1000638	F-HEMBB1000638	1124	R-HEMBB1000638	6634	
	HEMBB1000643	F-HEMBB1000643	1125	R-HEMBB1000643	6635	
	HEMBB1000649	F-HEMBB1000649	1126	R-HEMBB1000649	6636	
10	HEMBB1000652	F-HEMBB1000652	1127	R-HEMBB1000652	6637	
,,	HEMBB1000665	F-HEMBB1000665	1128	R-HEMBB1000665	6638	
	HEMBB1000671	F-HEMBB1000671	1129	R-HEMBB1000671	6639	
	HEMBB1000673	F-HEMBB1000673	1130	R-HEMBB1000673	6640	
15	HEMBB1000684	F-HEMBB1000684	1130	R-HEMBB1000684	6641	
13	HEMBB1000693	F-HEMBB1000693				
			1132	R-nnnnnnnnnnnn	6642	
	HEMBB1000705	F-HEMBB1000705	1133	R-HEMBB1000705	6643	
••	HEMBB1000706	F-HEMBB1000706	1134	R-HEMBB1000706	6644	
20	HEMBB1000709	F-HEMBB1000709	1135	R-HEMBB1000709	6645	
	HEMBB1000725	F-HEMBB1000725	1136	R-HEMBB1000725	6646	
	HEMBB1000726	F-HEMBB1000726	1137	R-HEMBB1000726	6647	
	HEMBB1000738	F-HEMBB1000738	1138	R-HEMBB1000738	6648	
25	HEMBB1000749	F-HEMBB1000749	1139	R-HEMBB1000749	6649	
	HEMBB1000763	F-HEMBB1000763	1140	R-HEMBB1000763	6650	
	HEMBB1000770	F-HEMBB1000770	1141	R-HEMBB1000770	6651	
	HEMBB1000774	F-HEMBB1000774	1142			
30	HEMBB1000781	F-HEMBB1000781	1143	R-HEMBB1000781	6652	
	HEMBB1000789	F-HEMBB1000789	1144	R-HEMBB1000789	6653	
	HEMBB1000790	F-HEMBB1000790	1145	R-HEMBB1000790	6654	
	HEMBB1000794	F-HEMBB1000794	1146	R-HEMBB1000794	6655	
35	HEMBB1000807	F-HEMBB1000807	1147	R-HEMBB1000807	6656	
	HEMBB1000810	F-HEMBB1000810	1148	R-HEMBB1000810	6657	
	HEMBB1000821	F-HEMBB1000821	1149	R-HEMBB1000821	6658	
	HEMBB1000822	F-HEMBB1000822	1150	R-HEMBB1000822	6659	
40	HEMBB1000826	F-HEMBB1000826	1151	R-HEMBB1000826	6660	
	HEMBB1000827	F-HEMBB1000827	1152	R-HEMBB1000827	6661	
	HEMBB1000831	F-HEMBB1000831	1153	R-HEMBB1000831	6662	
	HEMBB1000835	F-HEMBB1000835	1154	R-HEMBB1000835	6663	
45	HEMBB1000840	F-HEMBB1000840	1155	R-HEMBB1000840	6664	
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	HEMBB1000852	F-HEMBB1000852	1157	R-HEMBB1000852	6666	
	HEMBB1000870	F-HEMBB1000870	1158	R-HEMBB1000870	6667	
50	HEMBB1000876	F-HEMBB1000876	1159	R-HEMBB1000876	6668	
50	HEMBB1000883	F-HEMBB1000883	1160	R-HEMBB1000883	6669	
	HEMBB1000887	F-HEMBB1000887	1161	R-HEMBB1000887	6670	
	HEMBB1000888	F-HEMBB1000888	1162	R-HEMBB1000888	6671	
	HEMBB1000890	F-HEMBB1000890	1163	R-HEMBB1000890	6672	
55	HEMBB1000893	F-HEMBB1000893	1164	R-HEMBB1000893	6673	
	HEMBB1000908	F-HEMBB1000908	1165	R-HEMBB1000908	6674	

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	HEMBB1000910	F-HEMBB1000910	1166	R-HEMBB1000910	6675
	HEMBB1000913	F-HEMBB1000913	1167	R-HEMBB1000913	6676
	HEMBB1000915	F-HEMBB1000915	1168	R-HEMBB1000915	6677
5	HEMBB1000917	F-HEMBB1000917	1169	R-HEMBB1000917	6678
•	HEMBB1000927	F-HEMBB1000927	1170	R-HEMBB1000927	6679
	HEMBB1000947	F-HEMBB1000947	1171	R-HEMBB1000947	6680
	HEMBB1000959	F-HEMBB1000959	1172	R-HEMBB1000959	6681
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10	HEMBB1000975	F-HEMBB1000975	1174	R-HEMBB1000975	6683
	HEMBB1000981	F-HEMBB1000981	1175	R-HEMBB1000981	6684
	HEMBB1000985	F-HEMBB1000985		R-HEMBB1000985	6685
	HEMBB1000991		1176		
15		F-HEMBB1000991	1177	R-HEMBB1000991	6686
	HEMBB1000996	F-HEMBB1000996	1178	R-HEMBB1000996	6687
	HEMBB1001004	F-HEMBB1001004,		R-HEMBB1001004	6688
	HEMBB1001008	F-HEMBB1001008	1180	R-HEMBB1001008	6689
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	HEMBB1001020	F-HEMBB1001020	1183	R-HEMBB1001020	6692
	HEMBB1001024	F-HEMBB1001024	1184	R-HEMBB1001024	6693
25	HEMBB1001037	F-HEMBB1001037	1185	R-HEMBB1001037	6694
	HEMBB1001047	F-HEMBB1001047	1186	R-HEMBB1001047	6695
	HEMBB1001051	F-HEMBB1001051	1187	R-HEMBB1001051	6696
	HEMBB1001056	F-HEMBB1001056	1188	R-HEMBB1001056	6697
30	HEMBB1001058	F-HEMBB1001058	1189	R-HEMBB1001058	6698
00	HEMBB1001060	F-HEMBB1001060	1190	R-HEMBB1001060	6699
	HEMBB1001063	F-HEMBB1001063	1191	R-HEMBB1001063	6700
	HEMBB1001068	F-HEMBB1001068	1192	R-HEMBB1001068	6701
0.5	HEMBB1001096	F-HEMBB1001096	1193	R-HEMBB1001096	6702
35	HEMBB1001102	F-HEMBB1001102	1194	R-HEMBB1001102	6703
	HEMBB1001105	F-HEMBB1001105	1195	R-HEMBB1001105	6704
	HEMBB1001112	F-HEMBB1001112	1196		
	HEMBB1001114	F-HEMBB1001114	1197	R-HEMBB1001114	6705
40	HEMBB1001117	F-HEMBB1001117	1198	R-HEMBB1001117	6706
	HEMBB1001119	F-HEMBB1001119	1199	R-HEMBB1001119	6707
	HEMBB1001126	F-HEMBB1001126	1200	R-HEMBB1001126	6708
	HEMBB1001133	F-HEMBB1001133	1201	R-HEMBB1001133	6709
45	HEMBB1001137	F-HEMBB1001137	1202	R-HEMBB1001137	6710
	HEMBB1001142	F-HEMBB1001142	1203	R-HEMBB1001142	6711
	HEMBB1001151	F-HEMBB1001151	1204	R-HEMBB1001151	6712
	HEMBB1001153	F-HEMBB1001153	1205	R-HEMBB1001153	6713
50	HEMBB1001169	F-HEMBB1001169	1206	R-HEMBB1001169	6714
	HEMBB1001175	F-HEMBB1001175	1207	R-nnnnnnnnnnn	6715
	HEMBB1001177	F-HEMBB1001177	1208	R-HEMBB1001177	6716
	HEMBB1001182	F-HEMBB1001182	1209	R-HEMBB1001182	6717
55	HEMBB1001199	F-HEMBB1001199	1210	R-HEMBB1001199	6718
	HEMBB1001208	F-HEMBB10011208	1211	R-HEMBB1001208	6719
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	HEMBB1001209	F-HEMBB1001209	1212	D_UEI/D01001001	6720
	HEMBB1001210	F-HEMBB1001210	1212	R-HEMBB1001209 R-HEMBB1001210	
	HEMBB1001218		1213		6721
_		F-HEMBB1001218		R-HEMBB1001218	6722
5	HEMBB1001221	F-HEMBB1001221	1215	R-HEMBB1001221	6723
	HEMBB1001234	F-HEMBB1001234	1216	R-HEMBB1001234	6724
	HEMBB1001242	F-HEMBB1001242	1217	R-HEMBB1001242	6725
	HEMBB1001249	F-HEMBB1001249	1218	R-HEMBB1001249	6726
10	HEMBB1001253	F-HEMBB1001253	1219	R-HEMBB1001253	6727
	HEMBB1001254	F-HEMBB1001254	1220	R-HEMBB1001254	6728
	HEMBB1001267	F-HEMBB1001267	1221	R-HEMBB1001267	6729
	HEMBB1001271	F-HEMBB1001271	1222	R-HEMBB1001271	6730
15	HEMBB1001282	F-HEMBB1001282	1223	R-HEMBB1001282	6731
	HEMBB1001288	F-HEMBB1001288	1224	R-HEMBB1001288	6732
	HEMBB1001289	F-HEMBB1001289,	1225	R-HEMBB1001289	6733
	HEMBB1001294	F-HEMBB1001294	1226	R-HEMBB1001294	6734
20	HEMBB1001302	F-HEMBB1001302	1227	R-HEMBB1001302	6735
	HEMBB1001304	F-HEMBB1001304	1228	R-HEMBB1001304	6736
	HEMBB1001314	F-HEMBB1001314	1229	R-HEMBB1001314	6737
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25	HEMBB1001317	F-HEMBB1001317	1231	R-HEMBB1001317	6739
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	HEMBB1001331	F-HEMBB1001331	1233	R-HEMBB1001331	6741
	HEMBB1001335	F-HEMBB1001335	1234	R-HEMBB1001335	6742
20	HEMBB1001337	F-HEMBB1001337	1235	R-HEMBB1001337	6743
30	HEMBB1001339	F-HEMBB1001339	1236	R-HEMBB1001339	6744
	HEMBB1001346	F-HEMBB1001346	1237	R-HEMBB1001346	6745
	HEMBB1001348	F-HEMBB1001348	1238	R-HEMBB1001348	6746
	HEMBB1001356	F-HEMBB1001356	1239	R-HEMBB1001356	6747
35	HEMBB1001364	F-HEMBB1001364	1240	R-HEMBB1001364	6748
	HEMBB1001366	F-HEMBB1001366	1241	R-HEMBB1001366	6749
	HEMBB1001367	F-HEMBB1001367	1242	R-HEMBB1001367	6750
	HEMBB1001369	F-HEMBB1001369	1243	R-HEMBB1001369	6751
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	HEMBB1001384	F-HEMBB1001384	1245	R-HEMBB1001384	6753
	HEMBB1001387	F-HEMBB1001387	1246	R-HEMBB1001387	6754
	HEMBB1001394	F-HEMBB1001394	1247	R-HEMBB1001394	6755
45	HEMBB1001410	F-HEMBB1001410	1248	R-HEMBB1001410	6756
	HEMBB1001424	F-HEMBB1001424	1249	R-HEMBB1001424	6757
	HEMBB1001426	F-HEMBB1001426	1250	R-HEMBB1001426	6758
	HEMBB1001429	F-HEMBB1001429	1251	R-HEMBB1001429	6759
50	HEMBB1001436	F-HEMBB1001436	1252	R-HEMBB1001436	6760
	HEMBB1001443	F-HEMBB1001443	1253	R-HEMBB1001443	6761
	HEMBB1001449	F-HEMBB1001449	1254	R-HEMBB1001449	6762
	HEMBB1001454	F-HEMBB1001454	1255	R-HEMBB1001454	6763
55	HEMBB1001458	F-HEMBB1001458	1256	R-HEMBB1001458	6764
	HEMBB1001463	F-HEMBB1001463	1257	R-HEMBB1001463	6765
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		HEMBB1001464	F-HEMBB1001464	1258	R-HEMBB1001464	6766
		HEMBB1001482	F-HEMBB1001482	1259	R-HEMBB1001482	6767
		HEMBB1001500	F-HEMBB1001500	1260	R-HEMBB1001500	6768
5		HEMBB1001521	F-HEMBB1001521	1261	R-HEMBB1001521	6769
		HEMBB1001527	F-HEMBB1001527	1262	R-HEMBB1001527	6770
		HEMBB1001531	F-HEMBB1001531	1263	R-HEMBB1001531	6771
		HEMBB1001535	F-HEMBB1001535	1264	R-HEMBB1001535	6772
10	1	HEMBB1001536	F-HEMBB1001536	1265	R-HEMBB1001536	6773
,,,		HEMBB1001537	F-HEMBB1001537	1266	R-HEMBB1001537	6774
		HEMBB1001555	F-HEMBB1001555	1267	R-HEMBB1001555	6775
		HEMBB1001562	F-HEMBB1001562	1268	R-HEMBB1001562	6776
		HEMBB1001564	F-HEMBB1001564	1269	R-HEMBB1001564	6777
15	•					67 7 8
		HEMBB1001565	F-HEMBB1001565	1270	R-HEMBB1001565	
		HEMBB1001585	F-HEMBB1001585	1271	R-HEMBB1001585	6779
		HEMBB1001586	F-HEMBB1001586	1272	R-HEMBB1001586	6780
20	1	HEMBB1001588	F-HEMBB1001588	1273	R-HEMBB1001588	6781
		HEMBB1001603	F-HEMBB1001603	1274	R-HEMBB1001603	6782
		HEMBB1001618	F-HEMBB1001618	1275	R-HEMBB1001618	6783
		HEMBB1001619	F-HEMBB1001619	1276	R-HEMBB1001619	6784
25		HEMBB1001630	F-HEMBB1001630	1277	R-HEMBB1001630	6785
		HEMBB1001635	F-HEMBB1001635	1278	R-HEMBB1001635	6786
		HEMBB1001637	F-HEMBB1001637	1279	R-HEMBB1001637	6787
		HEMBB1001641	F-HEMBB1001641	1280	R-HEMBB1001641	6788
30		HEMBB1001653	F-HEMBB1001653	1281	R-HEMBB1001653	6789
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		HEMBB1001668	F-HEMBB1001668	1283	R-HEMBB1001668	6791
		HEMBB1001673	F-HEMBB1001673	1284	R-HEMBB1001673	6792
35		HEMBB1001684	F-HEMBB1001684	1285	R-HEMBB1001684	6793
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		HEMBB1001695	F-HEMBB1001695	1287	R-HEMBB1001695	6795
		HEMBB1001704	F-HEMBB1001704	1288	R-HEMBB1001704	6796
40		HEMBB1001706	F-HEMBB1001706	1289	R-HEMBB1001706	6797
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		HEMBB1001717	F-HEMBB1001717	1291	R-HEMBB1001717	6799
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45		HEMBB1001747	F-HEMBB1001747	1294	R-HEMBB1001747	6802
		HEMBB1001749	F-HEMBB1001749	1295	R-HEMBB1001749	6803
		HEMBB1001753	F-HEMBB1001753	1296	R-HEMBB1001753	6804
		HEMBB1001756	F-HEMBB1001756	1297	R-HEMBB1001756	6805
50	HEMBB1001760	F-HEMBB1001760	1298	R-HEMBB1001760	6806	
		HEMBB1001762	F-HEMBB1001762	1299	R-HEMBB1001762	6807
		HEMBB1001785	F-HEMBB1001785	1300	R-HEMBB1001785	6808
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55	HEMBB1001802	F-HEMBB1001802	1302	R-HEMBB1001802	6810	
	HEMBB1001812	F-HEMBB1001812	1303	R-HEMBB1001812	6811	

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	HEMBB1001831	F-HEMBB1001831	1305	R-HEMBB1001831	6813
	HEMBB1001834	F-HEMBB1001834	1306	•	
5	HEMBB1001836	F-HEMBB1001836	1307	R-HEMBB1001836	6814
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	HEMBB1001863	F-HEMBB1001863	1310	R-HEMBB1001863	6817
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	HEMBB1001868	F-HEMBB1001868	1312	R-HEMBB1001868	6819
	HEMBB1001869	F-HEMBB1001869	1313	R-HEMBB1001869	6820
	HEMBB1001872	F-HEMBB1001872	1314	R-HEMBB1001872	6821
15	HEMBB1001874	F-HEMBB1001874	1315	R-HEMBB1001874	6822
,0	HEMBB1001875	F-HEMBB1001875	1316	R-HEMBB1001875	6823
	HEMBB1001880	F-HEMBB1001880,	1317	R-HEMBB1001880	6824
	HEMBB1001899	F-HEMBB1001899	1318	R-HEMBB1001899	6825
20	HEMBB1001905	F-HEMBB1001905	1319	R-HEMBB1001905	6826
20	HEMBB1001906	F-HEMBB1001906	1320	R-HEMBB1001906	6827
	HEMBB1001908	F-HEMBB1001908	1321	R-HEMBB1001908	6828
	HEMBB1001910	F-HEMBB1001910	1322	R-HEMBB1001910	6829
0.5	HEMBB1001911	F-HEMBB1001911	1323	R-HEMBB1001911	6830
25	HEMBB1001915	F-HEMBB1001915	1324	R-HEMBB1001915	6831
	HEMBB1001921	F-HEMBB1001921	1325	R-HEMBB1001921	6832
	HEMBB1001922	F-HEMBB1001922	1326	R-HEMBB1001922	6833
	HEMBB1001925	F-HEMBB1001925	1327	R-HEMBB1001925	6834
30	HEMBB1001930	F-HEMBB1001930	1328	R-HEMBB1001930	6835
	HEMBB1001944	F-HEMBB1001944	1329	R-HEMBB1001944	6836
	HEMBB1001945	F-HEMBB1001945	1330	R-HEMBB1001945	6837
	HEMBB1001947	F-HEMBB1001947	1331	R-HEMBB1001947	6838
35	HEMBB1001950	F-HEMBB1001950	1332	R-HEMBB1001950	6839
	HEMBB1001952	F-HEMBB1001952	1333	R-HEMBB1001952	6840
	HEMBB1001953	F-HEMBB1001953	1334	R-HEMBB1001953	6841
	HEMBB1001957	F-HEMBB1001957	1335	R-HEMBB1001957	6842
40	HEMBB1001962	F-HEMBB1001962	1336	R-HEMBB1001962	6843
	HEMBB1001967	F-HEMBB1001967	1337	R-HEMBB1001967	6844
	HEMBB1001973	F-HEMBB1001973	1338	R-HEMBB1001973	6845
	HEMBB1001983	F-HEMBB1001983	1339	R-HEMBB1001983	6846
45	HEMBB1001988	F-HEMBB1001988	1340	R-HEMBB1001988	6847
	HEMBB1001990	F-HEMBB1001990	1341	R-HEMBB1001990	6848
	HEMBB1001996	F-HEMBB1001996	1342	R-HEMBB1001996	6849
	HEMBB1001997	F-HEMBB1001997	1343	R-HEMBB1001997	6850
50	HEMBB1002002	F-HEMBB1002002	1344	R-HEMBB1002002	6851
	HEMBB1002005	F-HEMBB1002005	1345	R-HEMBB1002005	6852
	HEMBB1002009	F-HEMBB1002009	1346	R-HEMBB1002009	6853
	HEMBB1002015	F-HEMBB1002015	1347	R-HEMBB1002015	6854
55	HEMBB1002042	F-HEMBB1002042	1348	R-HEMBB1002042	6855
	HEMBB1002043	F-HEMBB1002043	1349	R-HEMBB1002043	6856

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	HEMBB1002044	F-HEMBB1002044	1350	R-HEMBB1002044	6857
	HEMBB1002045	F-HEMBB1002045	1351	R-HEMBB1002045	6858
	HEMBB1002049	F-HEMBB1002049	1352	R-HEMBB1002049	6859
5	HEMBB1002050	F-HEMBB1002050	1353	R-HEMBB1002050	6860
	HEMBB1002068	F-HEMBB1002068	1354	R-HEMBB1002068	6861
	HEMBB1002069	F-HEMBB1002069	1355	R-HEMBB1002069	6862
	HEMBB1002092	F-HEMBB1002092	1356	R-HEMBB1002092	6863
10	HEMBB1002094	F-HEMBB1002094	1357	R-HEMBB1002094	6864
	HEMBB1002115	F-HEMBB1002115	1358	R-HEMBB1002115	6865
	HEMBB1002134	F-HEMBB1002134	1359		
	HEMBB1002139	F-HEMBB1002139	1360	R-HEMBB1002139	6866
15	HEMBB1002142	F-HEMBB1002142	1361	R-HEMBB1002142	6867
10	HEMBB1002152	F-HEMBB1002152	1362	R-HEMBB1002152	6868
	HEMBB1002189	F-HEMBB1002189		R-HEMBB1002189	6869
	HEMBB1002190	F-HEMBB1002190	1364	R-HEMBB1002190	6870
20	HEMBB1002193	F-HEMBB1002193	1365	R-HEMBB1002193	6871
20	HEMBB1002217	F-HEMBB1002217	1366	R-HEMBB1002217	6872
	HEMBB1002218	F-HEMBB1002218	1367	R-HEMBB1002218	6873
	HEMBB1002232	F-HEMBB1002232	1368	R-HEMBB1002232	6874
0.5	HEMBB1002247	F-HEMBB1002247	1369	R-HEMBB1002247	6875
25	HEMBB1002249	F-HEMBB1002249	1370	R-HEMBB1002249	6876
	HEMBB1002254	F-HEMBB1002254	1371	R-HEMBB1002254	6877
	HEMBB1002255	F-HEMBB1002255	1372	R-HEMBB1002255	6878
	HEMBB1002266	F-HEMBB1002266	1373	R-HEMBB1002266	6879
30	HEMBB1002280	F-HEMBB1002280	1374	R-HEMBB1002280	6880
	HEMBB1002300	F-HEMBB1002300	1375	R-HEMBB1002300	6881
	HEMBB1002306	F-HEMBB1002306	1376	R-HEMBB1002306	6882
	HEMBB1002327	F-HEMBB1002327	1377	R-HEMBB1002327	6883
35	HEMBB1002329	F-HEMBB1002329	1378	R-HEMBB1002329	6884
	HEMBB1002340	F-HEMBB1002340	1379	R-HEMBB1002340	6885
	HEMBB1002342	F-HEMBB1002342	1380	R-HEMBB1002342	6886
	HEMBB1002358	F-HEMBB1002358	1381	R-HEMBB1002358	6887
40	HEMBB1002359	F-HEMBB1002359	1382	R-HEMBB1002359	6888
	HEMBB1002364	F-HEMBB1002364	1383	R-HEMBB1002364	6889
	HEMBB1002371	F-HEMBB1002371	1384	R-HEMBB1002371	6890
	HEMBB1002381	F-HEMBB1002381	1385	R-HEMBB1002381	6891
45	HEMBB1002383	F-HEMBB1002383	1386	R-HEMBB1002383	6892
	HEMBB1002387	F-HEMBB1002387	1387	R-HEMBB1002387	6893
	HEMBB1002409	F-HEMBB1002409	1388		
	HEMBB1002415	F-HEMBB1002415	1389	R-HEMBB1002415	6894
50	HEMBB1002425	F-HEMBB1002425	1390	R-HEMBB1002425	6895
	HEMBB1002442	F-HEMBB1002442	1391	R-HEMBB1002442	6896
	HEMBB1002453	F-HEMBB1002453	1392	R-HEMBB1002453	6897
	HEMBB1002457	F-HEMBB1002457	1393	R-HEMBB1002457	6898
55	HEMBB1002458	F-HEMBB1002458		R-HEMBB1002458	6899
	HEMBB1002477	F-HEMBB1002477	1395	R-HEMBB1002477	6900

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	HEMBB1002489	F-HEMBB1002489	1396	R-HEMBB1002489	6901
	HEMBB1002492	F-HEMBB1002492	1397	R-HEMBB1002492	6902
	HEMBB1002495	F-HEMBB1002495	1398	R-HEMBB1002495	6903
5	HEMBB1002502	F-HEMBB1002502	1399	R-HEMBB1002502	6904
	HEMBB1002509	F-HEMBB1002509	1400	R-HEMBB1002509	6905
	HEMBB1002510	F-HEMBB1002510	1401	R-HEMBB1002510	6906
	HEMBB1002520	F-HEMBB1002520	1402	R-HEMBB1002520	6907
10	HEMBB1002522	F-HEMBB1002522	1403	R-HEMBB1002522	6908
	HEMBB1002531	F-HEMBB1002531	1404	R-HEMBB1002531	6909
	HEMBB1002534	F-HEMBB1002534	1405	R-HEMBB1002534	6910
	HEMBB1002545	F-HEMBB1002545	1406	R-HEMBB1002545	6911
15	HEMBB1002550	F-HEMBB1002550	1407	R-HEMBB1002550	6912
15	HEMBB1002556	F-HEMBB1002556	1408	R-HEMBB1002556	6913
	HEMBB1002579	F-HEMBB1002579	1409	R-HEMBB1002579	6914
	HEMBB1002513	F-HEMBB1002582	1410	R-HEMBB1002582	6915
	HEMBB1002590	F-HEMBB1002590	1411	R-HEMBB1002590	
20	HEMBB1002596	F-HEMBB1002596		R-HEMBB1002596	6916 6017
	HEMBB1002590	F-HEMBB1002600	1412		6917
	HEMBB1002601	F-HEMBB1002601	1413	R-HEMBB1002600	6918
	HEMBB1002603	F-HEMBB1002603	1414	R-HEMBB1002601	6919 6920
25	HEMBB1002607		1415	R-HEMBB1002603	
	HEMBB1002610	F-HEMBB1002607 F-HEMBB1002610	1416	R-HEMBB1002607	6921
	HEMBB1002613		1417	R-HEMBB1002610	6922
		F-HEMBB1002613 F-HEMBB1002614	1418	R-HEMBB1002613	6923
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		F-HEMBB1002617	1420	R-HEMBB1002617	6925
	HEMBB1002623	F-HEMBB1002623	1421	R-HEMBB1002623	6926
	HEMBB1002635	F-HEMBB1002635	1422	R-HEMBB1002635	6927
35	HEMBB1002664	F-HEMBB1002664	1423	R-HEMBB1002664	6928
	HEMBB1002677	F-HEMBB1002677	1424	R-HEMBB1002677	6929
	HEMBB1002683	F-HEMBB1002683	1425	R-HEMBB1002683	6930
	HEMBB1002684	F-HEMBB1002684	1426	R-HEMBB1002684	6931
40	HEMBB1002686	F-HEMBB1002686	1427	R-HEMBB1002686	6932
,•	HEMBB1002692 HEMBB1002697	F-HEMBB1002692	1428	R-HEMBB1002692	6933
		F-HEMBB1002697	1429	R-HEMBB1002697	6934
	HEMBB1002699	F-HEMBB1002699	1430	R-HEMBB1002699	6935
45	HEMBB1002702	F-HEMBB1002702	1431	R-HEMBB1002702	6936
40	HEMBB1002705	F-HEMBB1002705	1432	R-HEMBB1002705	6937
	HEMBB1002712	F-HEMBB1002712	1433	R-HEMBB1002712	6938
	MAMMA1000009	F-MAMMA1000009	1434	R-MAMMA1000009	6939
	MAMMA1000019	F-MAMMA1000019	1435	R-MAMMA1000019	6940
50	MAMMA1000020	F-MAMMA1000020	1436	R-MAMMA1000020	6941
	MAMMA1000025	F-MAMMA1000025	1437	R-MAMMA1000025	6942
	MAMMA1000043	F-MAMMA1000043	1438	R-MAMMA1000043	6943
	MAMMA1000045	F-MAMMA1000045	1439	R-MAMMA1000045	6944
55	MAMMA1000055	F-MAMMA1000055	1440	R-MAMMA1000055	6945
	MAMMA1000057	F-MAMMA1000057	1441	R-MAMMA1000057	6946

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	MAMMA1000069	F-MAMMA1000069	1442	R-MAMMA1000069	6947
	MAMMA1000084	F-MAMMA1000084	1443	R-MAMMA1000084	6948
	MAMMA1000085	F-MAMMA1000085	1444	R-MAMMA1000085	6949
5	MAMMA1000092	F-MAMMA1000092	1445	R-MAMMA1000092	6950
	MAMMA1000103	F-MAMMA1000103	1446	R-MAMMA1000103	6951
	MAMMA1000117	F-MAMMA1000117	1447	R-MAMMA1000117	6952
	MAMMA1000129	F-MAMMA1000129	1448	R-MAMMA1000129	6953
10	MAMMA1000133	F-MAMMA1000133	1449	R-MAMMA1000133	6954
10	MAMMA1000134	F-MAMMA1000134	1450	R-MAMMA1000134	6955
	MAMMA1000139	F-MAMMA1000139	1451	R-MAMMA1000139	6956
	MAMMA1000143	F-MAMMA1000143	1452	R-MAMMA1000143	6957
45	MAMMA1000155	F-MAMMA1000155	1453	R-MAMMA1000155	6958
15	MAMMA1000163	F-MAMMA1000163	1454	R-MAMMA1000163	6959
	MAMMA1000171	F-MAMMA1000171,	1455	R-MAMMA1000171	6960
	MAMMA1000171	F-MAMMA1000171	1456	R-MAMMA1000171	6961
	MAMMA1000175	F-MAMMA1000175	1457	R-MAMMA1000175	6962
20	MAMMA1000173	F-MAMMA1000173	1458	R-MAMMA1000183	6963
	MAMMA1000198	F-MAMMA1000183	1459	R-MAMMA1000198	6964
	MAMMA1000198	F-MAMMA1000221	1460	R-MAMMA1000138	6965
	MAMMA1000227	F-MAMMA1000227	1461	R-MAMMA1000227	6966
25	MAMMA1000227	F-MAMMA1000221	1462	R-MAMMA1000221	6967
	MAMMA1000241	F-MAMMA1000241	1463	R-MAMMA1000251	6968
	MAMMA1000251	F-MAMMA1000251	1464	R-MAMMA1000254	6969
	MAMMA1000254	F-MAMMA1000257	1465	R-MAMMA1000257	6970
30	MAMMA1000257	F-MAMMA1000257	1466	R-MAMMA1000264	6971
	MAMMA1000264	F-MAMMA1000266	1467	R-MAMMA1000266	6972
	MAMMA1000270	F-MAMMA1000270	1468	R-MAMMA1000270	6973
	MAMMA1000270	F-MAMMA1000277	1469	R-MAMMA1000277	6974
35	MAMMA1000277	F-MAMMA1000277	1470	R-MAMMA1000278	6975
	MAMMA1000279	F-MAMMA1000279	1471	R-MAMMA1000279	6976
	MAMMA1000279	F-MAMMA1000279	1472	R-MAMMA1000284	6977
	MAMMA1000284	F-MAMMA1000287	1473	R-MAMMA1000287	6978
40	MAMMA1000287	F-MAMMA1000302	1474	R-MAMMA1000302	6979
	MAMMA1000302	F-MAMMA1000307	1475	R-MAMMA1000307	6980
	MAMMA1000307	F-MAMMA1000309	1476	R-MAMMA1000309	6981
	MAMMA1000309	F-MAMMA1000312	1477	R-MAMMA1000312	6982
45	MAMMA1000312	F-MAMMA1000313	1478	R-MAMMA1000313	6983
40	MAMMA1000313	F-MAMMA1000313	1479	R-MAMMA1000331	6984
	MAMMA1000331	F-MAMMA1000331	1480	R-MAMMA1000339	6985
	MAMMA1000340	F-MAMMA1000340	1481	R-MAMMA1000340	6986
50	MAMMA1000340	F-MAMMA1000340	1482	R-MAMMA1000348	6987
50	MAMMA1000348 MAMMA1000356	F-MAMMA1000348	1483	R-MAMMA1000348	6988
	MAMMA1000356	F-MAMMA1000360	1484	R-MAMMA1000360	6989
		F-MAMMA1000360	1485	R-MAMMA1000361	6990
	MAMMA1000361	F-MAMMA1000372	1486	R-MAMMA1000372	6991
55	MAMMA1000372 MAMMA1000385	F-MAMMA1000372	1487	R-MAMMA1000385	6992
	MAMMA1000385	L_WVmmv1000999	1401	W. WANTAMAN V	0334

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	. MAMMA1000388	F-MAMMA1000388	1488	R-MAMMA1000388	6993
	MAMMA1000395	F-MAMMA1000395	1489	R-MAMMA1000395	6994
	MAMMA1000402	F-MAMMA1000402	1490	R-MAMMA1000402	6995
5	MAMMA1000410	F-MAMMA1000410	1491	R-MAMMA1000410	6996
	MAMMA1000413	F-MAMMA1000413	1492	R-MAMMA1000413	6997
	MAMMA1000414	F-MAMMA1000414	1493	R-MAMMA1000414	6998
	MAMMA1000416	F-MAMMA1000416	1494	R-MAMMA1000416	6999
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	MAMMA1000422	F-MAMMA1000422	1496	R-MAMMA1000422	7001
	MAMMA1000423	F-MAMMA1000423	1497	R-MAMMA1000423	7002
	MAMMA1000424	F-MAMMA1000424	1498	R-MAMMA1000424	7003
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	MAMMA1000431	F-MAMMA1000431	1500	R-MAMMA1000431	7005
	MAMMA1000444	F-MAMMA1000444,	1501	R-MAMMA1000444	7006
	MAMMA1000446	F-MAMMA1000446	1502	R-MAMMA1000446	7007
20	MAMMA1000458	F-MAMMA1000458	1503	R-MAMMA1000458	7008
	MAMMA1000468	F-MAMMA1000468	1504	R-MAMMA1000468	7009
	MAMMA1000472	F-MAMMA1000472	1505	R-MAMMA1000472	7010
	MAMMA1000478	F-MAMMA1000478	1506	R-MAMMA1000478	7011
25	MAMMA1000483	F-MAMMA1000483	1507	R-MAMMA1000483	7012
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	MAMMA1000500	F-MAMMA1000500	1509	R-MAMMA1000500	7014
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30	MAMMA1000522	F-MAMMA1000522	1512	R-MAMMA1000522	7017
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35	MAMMA1000565	F-MAMMA1000565	1515	R-MAMMA1000565	7019
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	MAMMA1000583	F-MAMMA1000583	1518	R-MAMMA1000583	7022
40	MAMMA1000585	F-MAMMA1000585	1519	R-MAMMA1000585	7023
40	MAMMA1000594	F-MAMMA1000594	1520	R-MAMMA1000594	7024
	MAMMA1000597	F-MAMMA1000597	1521	R-MAMMA1000597	7025
	MAMMA1000605	F-MAMMA1000605	1522	R-MAMMA1000605	7026
45	MAMMA1000612	F-MAMMA1000612	1523	R-MAMMA1000612	7027
45	MAMMA1000616	F-MAMMA1000616	1524	R-MAMMA1000616	7028
	MAMMA1000621	F-MAMMA1000621	1525	R-MAMMA1000621	7029
	MAMMA1000623	F-MAMMA1000623	1526	R-MAMMA1000623	7030
	MAMMA1000625	F-MAMMA1000625	1527	R-MAMMA1000625	7031
50	MAMMA1000643	F-MAMMA1000643	1528	R-MAMMA1000643	7032
	MAMMA1000664	F-MAMMA1000664	1529	R-MAMMA1000664	7033
	MAMMA1000669	F-MAMMA1000669	1530	R-MAMMA1000669	7034
	MAMMA1000670	F-MAMMA1000670	1531	R-MAMMA1000670	7035
55	MAMMA1000672	F-MAMMA1000672	1532	R-MAMMA1000672	7036
	MAMMA1000684	F-MAMMA1000684	1533	R-MAMMA1000684	7037
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	MAMMA1000696	F-MAMMA1000696	1534	R-MAMMA1000696	7038
	MAMMA1000707	F-MAMMA1000707	1535	R-MAMMA1000707	7039
	MAMMA1000713	F-MAMMA1000713	1536	R-MAMMA1000713	7040
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	MAMMA1000718	F-MAMMA1000718	1538	R-MAMMA1000718	7042
	MAMMA1000720	F-MAMMA1000720	1539	R-MAMMA1000720	7043
	MAMMA1000723	F-MAMMA1000723	1540	R-MAMMA1000723	7044
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		F-MAMMA1000733		R-MAMMA1000732	7040
	MAMMA1000733		1543		
45	MAMMA1000734	F-MAMMA1000734	1544	R-MAMMA1000734	7048
15	MAMMA1000738	F-MAMMA1000738	1545	R-MAMMA1000738	7049
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	MAMMA1000746	F-MAMMA1000746	1547	R-MAMMA1000746	7051
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20	MAMMA1000760	F-MAMMA1000760	1549	R-MAMMA1000760	7053
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	MAMMA1000776	F-MAMMA1000776	1552	R-MAMMA1000776	7056
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	MAMMA1000802	F-MAMMA1000802	1556	R-MAMMA1000802	7060
30	MAMMA1000824	F-MAMMA1000824	1557		
••	MAMMA1000831	F-MAMMA1000831	1558	R-MAMMA1000831	7061
	MAMMA1000839	F-MAMMA1000839	1559	R-MAMMA1000839	7062
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35	MAMMA1000842	F-MAMMA1000842	1561	R-MAMMA1000842	7064
33	MAMMA1000843	F-MAMMA1000843	1562	R-MAMMA1000843	7065
	MAMMA1000845	F-MAMMA1000845	1563	R-MAMMA1000845	7066
	MAMMA1000851	F-MAMMA1000851	1564	R-MAMMA1000851	7067
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40	MAMMA 1000856	F-MAMMA1000856	1566	R-MAMMA1000856	7069
	MAMMA 1000859	F-MAMMA1000859	1567	•	
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45	MAMMA1000865	F-MAMMA1000865	1570	R-MAMMA1000865	7072
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	MAMMA1000876	F-MAMMA1000876	1573	R-MAMMA1000876	7075
50	MAMMA1000877	F-MAMMA1000877	1574	R-MAMMA1000877	7076
	MAMMA1000880	F-MAMMA1000880	1575	R-MAMMA1000880	7077
	MAMMA1000883	F-MAMMA1000883	1576	R-MAMMA1000883	7078
	MAMMA1000897	F-MAMMA1000897	1577	R-MAMMA1000897	7079
55	MAMMA1000905	F-MAMMA1000905	1578	R-MAMMA1000905	7080
	MAMMA1000905	F-MAMMA1000906	1579	R-MAMMA1000906	7081
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	MAMMA1000914	F-MAMMA1000914	1581	R-MAMMA1000914	7083
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	MAMMA1000941	F-MAMMA1000941	1585	R-MAMMA1000941	7087
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		F-MAMMA1000957	1589	R-MAMMA1000957	7091
	MAMMA1000962	F-MAMMA1000962	1590	R-MAMMA1000962	7092
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	MAMMA1000975	F-MAMMA1000975	1592	R-MAMMA1000975	7094
	MAMMA1000979	F-MAMMA1000979,	1593	R-MAMMA1000979	7095
	MAMMA1000987	F-MAMMA1000987	1594	R-MAMMA1000987	7096
20	MAMMA1000998	F-MAMMA1000998	1595	R-MAMMA1000998	7097
	MAMMA1001003	F-MAMMA1001003	1596	R-MAMMA1001003	7098
	MAMMA1001008	F-MAMMA1001008	1597	R-MAMMA1001008	7099
	MAMMA1001021	F-MAMMA1001021	1598	R-MAMMA1001021	7100
25	MAMMA1001024	F-MAMMA1001024	1599	R-MAMMA1001024	7101
	MAMMA1001030	F-MAMMA1001030	1600	R-MAMMA1001030	7102
	MAMMA1001035	F-MAMMA1001035	1601	R-MAMMA1001035	7103
	MAMMA1001038	F-MAMMA1001038	1602	R-MAMMA1001038	7104
30	MAMMA1001041	F-MAMMA1001041	1603	R-nnnnnnnnnnn	7105
30	MAMMA1001050	F-MAMMA1001050	1604	R-MAMMA1001050	7106
	MAMMA1001059	F-MAMMA1001059	1605	R-MAMMA1001059	7107
,	MAMMA1001067	F-MAMMA 1001067	1606	R-MAMMA1001067	7108
	MAMMA1001073	F-MAMMA1001073	1607	R-MAMMA1001073	7109
35	MAMMA1001074	F-MAMMA1001074	1608	R-MAMMA1001074	7110
	MAMMA1001075	F-MAMMA1001075	1609	R-MAMMA1001075	7111
	MAMMA1001078	F-MAMMA1001078	1610	R-MAMMA1001078	7112
	MAMMA1001080	F-MAMMA1001080	1611		
40	MAMMA1001082	F-MAMMA1001082	1612	R-MAMMA1001082	7113
	MAMMA1001091	F-MAMMA1001091	1613	R-MAMMA1001091	7114
	MAMMA1001092	F-MAMMA1001092	1614	R-MAMMA1001092	7115
•	MAMMA10.01105	F-MAMMA1001105	1615	R-MAMMA1001105	7116
45	MAMMA1001110	F-MAMMA1001110	1616	R-MAMMA1001110	7117
	MAMMA1001126	F-MAMMA1001126	1617	R-MAMMA1001126	7118
	MAMMA1001133	F-MAMMA1001133	1618	R-MAMMA1001133	7119
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50	MAMMA1001143	F-MAMMA1001143	1620	R-MAMMA1001143	7121
	MAMMA1001145	F-MAMMA1001145	1621	R-MAMMA1001145	7122
	MAMMA1001154	F-MAMMA1001154	1622	R-MAMMA1001154	7123
	MAMMA1001161	F-MAMMA1001161	1623	R-MAMMA1001161	7124
55	MAMMA1001162	F-MAMMA1001162	1624	R-MAMMA1001161	7125
•	MAMMA1001181	F-MAMMA1001181	1625	R-MAMMA1001181	7126
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	MAMMA1001186	F-MAMMA1001186	1626	R-MAMMA1001186	7127
	MAMMA1001191	F-MAMMA1001191	1627	R-MAMMA1001191	7128
	MAMMA1001198	F-MAMMA1001198	1628	R-MAMMA1001198	7129
5	MAMMA1001202	F-MAMMA1001202	1629	R-MAMMA1001202	7130
	MAMMA1001203	F-MAMMA1001203	1630	R-MAMMA1001203	7131
	MAMMA1001206	F-MAMMA1001206	1631	R-MAMMA1001206	7132
	MAMMA1001215	F-MAMMA1001215	1632	R-MAMMA1001215	7133
10	MAMMA1001220	F-MAMMA1001220	1633	R-MAMMA1001220	7134
	MAMMA1001222	F-MAMMA1001222	1634	R-MAMMA1001222	7135
	MAMMA1001243	F-MAMMA1001243	1635	R-MAMMA1001243	7136
	MAMMA1001244	F-MAMMA1001244	1636	R-MAMMA1001244	7137
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15	MAMMA1001256	F-MAMMA1001256	1638	R-MAMMA1001256	7139
	MAMMA1001259	F-MAMMA1001259	1639	R-MAMMA1001259	7140
	MAMMA1001260	F-MAMMA1001260	1640	R-MAMMA1001260	7141
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20	MAMMA1001271	F-MAMMA1001271	1642	R-MAMMA1001271	7143
	MAMMA1001274	F-MAMMA1001274	1643	R-MAMMA1001274	7144
	MAMMA1001280	F-MAMMA1001280	1644	R-MAMMA1001280	7145
	MAMMA1001292	F-MAMMA1001292	1645	R-MAMMA1001292	7146
25	MAMMA1001296	F-MAMMA1001296	1646	R-MAMMA1001296	7147
	MAMMA1001298	F-MAMMA1001298	1647	R-MAMMA1001298	7148
	MAMMA1001305	F-MAMMA1001305	1648	R-MAMMA1001305	7149
	MAMMA1001322	F-MAMMA1001322	1649	R-MAMMA1001322	7150
30	MAMMA1001324	F-MAMMA1001324	1650	R-MAMMA1001324	7151
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	MAMMA1001341	F-MAMMA1001341	1652	R-MAMMA1001341	7153
	MAMMA1001343	F-MAMMA1001343	1653	R-MAMMA1001343	7154
35	MAMMA1001346	F-MAMMA1001346	1654	R-MAMMA1001346	7155
	MAMMA1001383	F-MAMMA1001383	1655	R-MAMMA1001383	7156
	MAMMA1001388	F-MAMMA1001388	1656	R-MAMMA1001388	7157
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40	MAMMA1001408	F-MAMMA1001408	1658	R-MAMMA1001408	7159
	MAMMA1001411	F-MAMMA1001411	1659	R-MAMMA1001411	7160
	MAMMA1001419	F-MAMMA1001419	1660	R-MAMMA1001419	7161
	MAMMA1001420	F-MAMMA1001420	1661	R-MAMMA1001420	7162
45	MAMMA1001435	F-MAMMA1001435	1662	R-MAMMA1001435	7163
	MAMMA1001442	F-MAMMA1001442	1663	R-MAMMA1001442	7164
	MAMMA1001446	F-MAMMA1001446	1664	R-MAMMA1001446	7165
	MAMMA1001452	F-MAMMA1001452	1665	R-MAMMA1001452	7166
50	MAMMA1001465	F-MAMMA1001465	1666	R-MAMMA1001465	7167
30	MAMMA1001476	F-MAMMA1001476	1667	R-MAMMA1001476	7168
	MAMMA1001487	F-MAMMA1001487	1668	R-MAMMA1001487	7169
	MAMMA1001501	F-MAMMA1001501	1669	R-MAMMA1001501	7170
	MAMMA1001502	F-MAMMA1001502	1670	R-MAMMA1001502	7171
55	MAMMA1001510	F-MAMMA1001510	1671	R-MAMMA1001510	7172

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	MAMMA1001522	F-MAMMA1001522	1672	R-MAMMA1001522	7173
	MAMMA1001547	F-MAMMA1001547	1673	R-MAMMA1001547	7174
	MAMMA1001551	F-MAMMA1001551	1674	R-MAMMA1001551	7175
5	MAMMA1001575	F-MAMMA1001575	1675	R-MAMMA1001575	7176
	MAMMA1001576	F-MAMMA1001576	1676	R-MAMMA1001576	7177
	MAMMA1001590	F-MAMMA1001590	1677	R-MAMMA1001590	7178
	MAMMA1001600	F-MAMMA1001600	1678	R-MAMMA1001600	7179
10	MAMMA1001604	F-MAMMA1001604	1679	R-MAMMA1001604	7180
,,,	MAMMA1001606	F-MAMMA1001606	1680	R-MAMMA1001606	7181
	MAMMA1001620	F-MAMMA1001620	1681	R-MAMMA1001620	7182
	MAMMA1001627	F-MAMMA1001627	1682	R-MAMMA1001627	7183
15	MAMMA1001630	F-MAMMA1001630	1683	R-MAMMA1001630	7184
15	MAMMA1001633	F-MAMMA1001633	1684	R-MAMMA1001633	7185
	MAMMA1001635	F-MAMMA1001635	1685	R-MAMMA1001635	7186
	MAMMA1001649	F-MAMMA1001649	1686	R-MAMMA1001649	7187
	MAMMA1001654	F-MAMMA1001654	1687	It in the second	
20	MAMMA1001663	F-MAMMA1001663	1688	R-MAMMA1001663	7188
	MAMMA1001670	F-MAMMA1001670	1689	R-MAMMA1001670	7189
	MAMMA1001671	F-MAMMA1001671	1690	R-MAMMA1001671	7190
	MAMMA1001679	F-MAMMA1001679	1691	R-MAMMA1001679	7191
25	MAMMA1001683	F-MAMMA1001683	1692	R-MAMMA1001683	7192
	MAMMA1001686	F-MAMMA1001686	1693	R-MAMMA1001686	7193
	MAMMA1001692	F-MAMMA1001692	1694	R-MAMMA1001692	7194
	MAMMA1001711	F-MAMMA1001711	1695	R-MAMMA1001711	7195
30	MAMMA1001711	F-MAMMA1001715	1696	R-MAMMA1001715	7196
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	MAMMA1001735	F-MAMMA1001735	1698	R-MAMMA1001735	7198
	MAMMA1001740	F-MAMMA1001740	1699	R-MAMMA1001740	7199
35	MAMMA1001743	F-MAMMA1001743	1700	R-MAMMA1001743	7200
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	MAMMA1001815		1717	R-MAMMA1001815	7217

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45	MAMMA1001868	F-MAMMA1001868			7229
15	MAMMA1001874		1729	R-nnnnnnnnnnnn	7230
		F-MAMMA1001874	1730	R-MAMMA1001874	
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	MAMMA1001931	F-MAMMA1001931	1736	R-MAMMA1001931	7236
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40	MAMMA1002056	F-MAMMA1002056	1749	R-MAMMA1002056	7249
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	MAMMA1002078	F-MAMMA1002078	1752	R-MAMMA1002078	7252
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	MAMMA1002118	F-MAMMA1002118	1757	R-MAMMA1002118	7257
50	MAMMA1002125	F-MAMMA1002125	1758	R-MAMMA1002125	7258
	MAMMA1002132	F-MAMMA1002132	1759	R-MAMMA1002132	7259
	MAMMA1002140	F-MAMMA1002140	1760	R-MAMMA1002140	7260
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55	MAMMA1002145	F-MAMMA1002145	1762	R-MAMMA1002145	7262
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	MAMMA1002158	F-MAMMA1002158	1766	R-MAMMA1002158	7266
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	MAMMA1002298	F-MAMMA1002298	1786	R-MAMMA1002299	7285 7286
	MAMMA1002299	F-MAMMA1002299		R-MAMMA1002299 R-MAMMA1002308	7287
30	MAMMA1002308	F-MAMMA1002308	1787 1788	R-MAMMA1002310	7288
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	MAMMA1002311	F-MAMMA1002311 F-MAMMA1002312	1789 1790	R-MAMMA1002311 R-MAMMA1002312	7299 7290
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	MAMMA1002355	F-MAMMA1002355	1802	R-MAMMA1002355	7302
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	MAMMA1002359	F-MAMMA1002359	1804	R-MAMMA1002359	7304
50	MAMMA1002359	F-MAMMA1002360	1805	R-MAMMA1002360	7305
	MAMMA1002361	F-MAMMA1002361 F-MAMMA1002362	1806 1807	R-MAMMA1002361 R-MAMMA1002362	7306 7307
	MAMMA1002362	F-MAMMA1002380	1808	R-MAMMA1002380	7308
55	MAMMA1002380			R-MAMMA1002384	7309
	MAMMA1002384	F-MAMMA1002384	1809	л- мл ишл10023 0 4	1303

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	MAMMA1002385	F-MAMMA1002385	1810	R-MAMMA1002385	7310
	MAMMA1002392	F-MAMMA1002392	1811	R-MAMMA1002392	7311
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•	MAMMA1002417	F-MAMMA1002417	1814	R-MAMMA1002417	7314
	MAMMA1002417	F-MAMMA1002427	1815	R-MAMMA1002427	7315
	MAMMA1002428	F-MAMMA1002428	1816	R-MAMMA1002428	7316
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	MAMMA1002454	F-MAMMA1002454	1819	R-MAMMA1002454	7319
	MAMMA1002461	F-MAMMA1002461	1820	R-MAMMA1002461	7320
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	MAMMA1002475	F-MAMMA1002475	1822	R-MAMMA1002475	7322
•	MAMMA1002480	F-MAMMA1002480,	1823	R-MAMMA1002480	7323
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20	MAMMA1002494	F-MAMMA1002494	1825	R-MAMMA1002494	7325
20	MAMMA1002498	F-MAMMA1002498	1826	R-MAMMA1002498	7326
	MAMMA1002524	F-MAMMA1002524	1827	R-MAMMA1002524	7327
	MAMMA1002530	F-MAMMA1002530	1828	R-MAMMA1002530	7328
	MAMMA1002545	F-MAMMA1002545	1829	R-MAMMA1002545	7329
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	MAMMA1002556	F-MAMMA1002556	1831	R-MAMMA1002556	7331
•	MAMMA1002566	F-MAMMA1002566	1832	R-MAMMA1002566	7332
	MAMMA1002571	F-MAMMA1002571	1833	R-MAMMA1002571	7333
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	MAMMA1002585	F-MAMMA1002585	1835	R-MAMMA1002585	7335
	MAMMA1002590	F-MAMMA1002590	1836	R-MAMMA1002590	7336
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43	MAMMA1002629	F-MAMMA 1002629	1847	R-MAMMA1002629	7347
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50	MAMMA1002650	F-MAMMA1002650	1851	R-MAMMA1002650	7351
	MAMMA1002655	F-MAMMA1002655	1852	R-MAMMA1002655	7352
	MAMMA1002662	F-MAMMA1002662	1853	R-MAMMA1002662	7353
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	MAMMA1002684	F-MAMMA1002684	1857	R-MAMMA1002684	7357
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	MAMMA1002728	F-MAMMA1002728	1866	R-MAMMA1002728	7366
	MAMMA1002744	F-MAMMA1002744	1867	R-MAMMA1002744	7367
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	MAMMA1002748	F-MAMMA1002748,		R-MAMMA1002748	
	MAMMA1002754	F-MAMMA1002754	1870	R-MAMMA1002754	7370
20	MAMMA1002758	F-MAMMA1002758	1871	R-MAMMA1002758	7371
	MAMMA1002764	F-MAMMA1002764	1872	R-MAMMA1002764	7372
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25	MAMMA1002775	F-MAMMA1002775	1875	D 141044400500	7077
	MAMMA1002780	F-MAMMA1002780	1876	R-MAMMA1002780	7375
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	MAMMA1002833	F-MAMMA1002833	1882	R-MAMMA1002833	7381
35	MAMMA1002835	F-MAMMA1002835	1883	R-MAMMA1002835	7382
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	MAMMA1002895	F-MAMMA1002895	1898	R-MAMMA1002895	7396
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55	MAMMA1002909	F-MAMMA1002909	1900	R-MAMMA1002909	7398
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	MAMMA1002941	F-MAMMA1002941	1904	R-MAMMA1002941	7401
5	MAMMA1002947	F-MAMMA1002947	1905	R-MAMMA1002947	7402
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15	MAMMA1003004	F-MAMMA1003004	1913	R-MAMMA1003004	7410
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	MAMMA1003011	F-MAMMA1003011,	1915	R-MAMMA1003011	7412
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	MAMMA1003026	F-MAMMA1003026	1919	R-MAMMA1003026	7415 ·
	MAMMA1003031	F-MAMMA1003031	1920	R-MAMMA1003031	7416
25	MAMMA1003035	F-MAMMA1003035	1921	R-MAMMA1003035	7417
	MAMMA1003039	F-MAMMA1003039	1922	R-MAMMA1003039	7418
	MAMMA1003040	F-MAMMA1003040	1923	R-MAMMA1003040	7419
	MAMMA1003044	F-MAMMA1003044	1924	R-MAMMA1003044	7420
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	MAMMA1003055	F-MAMMA1003055	1927	R-MAMMA1003055	7423
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35	MAMMA1003057	F-MAMMA1003057	1929	R-MAMMA1003057	7425
	MAMMA1003066	F-MAMMA1003066	1930	R-MAMMA1003066	7426
	MAMMA1003089	F-MAMMA1003089	1931	R-MAMMA1003089	7427
	MAMMA1003099	F-MAMMA1003099	1932	R-MAMMA1003099	7428
40	MAMMA1003104	F-MAMMA1003104	1933	R-MAMMA1003104	7429
	MAMMA1003113	F-MAMMA1003113	1934	R-MAMMA1003113	7430
	MAMMA1003127	F-MAMMA1003127	1935	R-MAMMA1003127	7431
	MAMMA1003135	F-MAMMA1003135	1936	R-MAMMA1003135	7432
45	MAMMA1003140	F-MAMMA1003140	1937	R-MAMMA1003140	7433
	MAMMA1003146	F-MAMMA1003146	1938	R-MAMMA1003146	7434
	MAMMA1003150	F-MAMMA1003150	1939	R-nnnnnnnnnnnn	7435
	MAMMA1003166	F-MAMMA1003166	1940	R-MAMMA1003166	7436
50	NT2RM1000001	F-NT2RM1000001	1941		
	NT2RM1000018	F-NT2RM1000018	1942		
	NT2RM1000032	F-NT2RM1000032	1943		
	NT2RM1000035	F-NT2RM1000035	1944		
55	NT2RM1000037	F-NT2RM1000037	1945		
	NT2RM1000039	F-NT2RM1000039	1946		
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	NT2RM1000059	F-NT2RM1000059	1948
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33	NT2RM1000355	F-NT2RM1000355	1976
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-	NT2RM1000388	F-NT2RM1000388	1979
40	NT2RM1000394	F-NT2RM1000394	1980
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50	NT2RM1000563	F-NT2RM1000563	1988
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	NT2RM1000661	F-NT2RM1000661	1991
55	NT2RM1000666	F-NT2RM1000666	1992
	NT2RM1000669	F-NT2RM1000669	1993

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	NT2RM1000672	F-NT2RM1000672	1994		
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	NT2RM1000699	F-NT2RM1000699	1996		
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55	MADDI 1001102	T MTODMIOOTIVE	2000		

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55	NT2RM2000594		2085
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55	NT2RM2001370	F-NT2RM2001370	2131

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35	NT2RM2001695	F-NT2RM2001695	2160
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55	NT2RM2001784	F-NT2RM2001784	2177
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	NT2RM4000104	F-NT2RM4000104	2223	R-NT2RM4000104	7445

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	NT2RM4000425	F-NT2RM4000425	2254	R-NT2RM4000425	7476
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	NT2RM4000700	F-NT2RM4000700	2276	R-nnnnnnnnnnn	7498
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25	NT2RP2004463	F-NT2RP2004463	2980	N 112M 2007420	0040
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30	NT2RP2005031	F-NT2RP2005031	3030	R-NT2RP2005031	8095
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70	NT2RP2005147	F-NT2RP2005147	3039	R-NT2RP2005147	8104
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	NT2RP2005675	F-NT2RP2005675	3097	R-NT2RP2005675	8160
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	NT2RP2006219	F-NT2RP2006219	3150	R-NT2RP2006219	8210
10	NT2RP2006237	F-NT2RP2006237	3151	R-NT2RP2006237	8211
	NT2RP2006238	F-NT2RP2006238	3152	R-NT2RP2006238	8212
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	NT2RP2006365	F-NT2RP2006365	3162	R-NT2RP2006365	8221
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25	NT2RP3000759	F-NT2RP3000759	3255	R-NT2RP3000759	8309
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	NT2RP3001392	F-NT2RP3001392	3324	R-NT2RP3001392	8376
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	NT2RP3001428	F-NT2RP3001428	3332	R-nnnnnnnnnnn	8384
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	NT2RP4002052	F-NT2RP4002052	3818	R-NT2RP4002052	8837
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40	NT2RP4002071	F-NT2RP4002071	3820	R-NT2RP4002071	8839
	NT2RP4002075	F-NT2RP4002075	3821	R-NT2RP4002075	8840
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	NT2RP5003477	F-NT2RP5003477	3831		
55	NT2RP5003492	F-NT2RP5003492	3832		
	NT2RP5003500	F-NT2RP5003500	3833		ļ

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	NT2RP5003506	F-NT2RP5003506	3834		I
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5	NT2RP5003524	F-NT2RP5003524	3837		I
	NT2RP5003534	F-NT2RP5003534	3838		I
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	OVARC1000013	F-0VARC1000013	3842	R-0VARC1000013	8851
	OVARC1000014	F-0VARC1000014	3843	R-0VARC1000014	8852
	OVARC1000017	F-0VARC1000017	3844	R-0VARC1000017	8853
15	OVARC1000035	F-0VARC1000035	3845	R-0VARC1000035	8854
10	OVARC1000058	F-0VARC1000058	3846	R-0VARC1000058	8855
	OVARC1000060	F-0VARC1000060,	3847	R-0VARC1000060	8856
	OVARC1000068	F-0VARC1000068	3848	R-0VARC1000068	8857
20	OVARC1000071	F-0VARC1000071	3849	R-0VARC1000071	8858
20	OVARC1000085	F-0VARC1000085	3850	R-0VARC1000085	8859
	OVARC1000087	F-0VARC1000087	3851	R-nnnnnnnnnnnn	8860
	OVARC1000091	F-0VARC1000091	3852	R-0VARC1000091	8861
0.5	OVARC1000092	F-0VARC1000092	3853	R-0VARC1000092	8862
25	OVARC1000106	F-0VARC1000106	3854	R-0VARC1000106	8863
	OVARC1000109	F-0VARC1000109	3855	** *··	
	OVARC1000113	F-0VARC1000113	3856	R-0VARC1000113	8864
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30	OVARC1000133	F-0VARC1000133	3858	R-0VARC1000133	8866
	OVARC1000139	F-0VARC1000139	3859		
	OVARC1000145	F-0VARC1000145	3860	R-OVARC1000145	8867
~*	OVARC1000148	F-0VARC1000148	3861	R-0VARC1000148	8868
35	OVARC1000151	F-0VARC1000151	3862	R-0VARC1000151	8869
	OVARC1000168	F-0VARC1000168	3863	R-0VARC1000168	8870
	OVARC1000191	F-0VARC1000191	3864	R-0VARC1000191	8871
	0VARC1000198	F-0VARC1000198	3865	R-0VARC1000198	8872
40	0VARC1000209	F-0VARC1000209	3866	R-0VARC1000209	8873
	OVARC1000212	F-0VARC1000212	3867	R-0VARC1000212	8874
	0VARC1000240	F-0VARC1000240	3868	R-0VARC1000240	8875
	OVARC1000241	F-0VARC1000241	3869	R-OVARC1000241	8876
45	0VARC1000288	F-OVARC1000288	3870	R-0VARC1000288	8877
	0VARC1000302	F-OVARC1000302	3871	R-0VARC1000302	8878
	OVARC1000304	F-0VARC1000304	3872	R-0VARC1000304	8879
	0VARC1000309	F-0VARC1000309	3873	R-0VARC1000309	8880
50	0VARC1000321	F-0VARC1000321	3874	R-0VARC1000321	8881
	OVARC1000326	F-0VARC1000326	3875	R-0VARC1000326	8882
	OVARC1000335	F-0VARC1000335	3876	R-0VARC1000335	8883
	0VARC1000347	F-0VARC1000347	3877	R-0VARC1000347	8884
55	0VARC1000384	F-0VARC1000384	3878	R-OVARC1000384	8885
	0VARC1000408	F-0VARC1000408	3879	R-0VARC1000408	8886

		EP 107	4 61 <i>1 P</i>	12	
	0VARC1000411	F-0VARC1000411	3880	R-0VARC1000411	8887
	OVARC1000414	F-0VARC1000414	3881	R-0VARC1000414	8888
	0VARC1000420	F-OVARC1000420	3882	R-OVARC1000420	8889
5	0VARC1000427	F-0VARC1000427	3883	R-0VARC1000427	8890
	OVARC1000431	F-0VARC1000431	3884	R-OVARC1000431	8891
	0VARC1000437	F-0VARC1000437	3885	R-OVARC1000437	8892
	0VARC1000440	F-0VARC1000440	3886	R-0VARC1000440	8893
10	0VARC1000442	F-0VARC1000442	3887	R-OVARC1000442	8894
,,	0VARC1000443	F-0VARC1000443	3888	R-0VARC1000443	8895
	OVARC1000461	F-0VARC1000461	3889	R-0VARC1000461	8896
	0VARC1000465	F-0VARC1000465	3890	R-0VARC1000465	8897
15	0VARC1000466	F-0VARC1000466	3891	R-0VARC1000466	8898
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	0VARC1000473	F-0VARC1000479,	3893	R-0VARC1000479	8900
	0VARC1000415	F-0VARC1000479,	3894	R-0VARC1000486	8901
	0VARC1000486	F-0VARC1000496	3895	R-OVARC1000496	8902
20	0VARC1000490	F-0VARC1000430	3896	R-0VARC1000490	8903
	0VARC1000520	F-0VARC1000526	3897	R-OVARC1000526	8904
	0VARC1000528	F-0VARC1000528	3898	R-OVARC1000528	8905
	0VARC1000533	F-0VARC1000543	3899	R-0VARC1000543	8906
25	0VARC1000545	F-0VARC1000556	3900	R-0VARC1000545	8907
	0VARC1000557	F-0VARC1000557	3901	R-0VARC1000557	8908
	0VARC1000557 0VARC1000564	F-0VARC1000557	3902	R-0VARC1000557	8909
	0VARC1000504 0VARC1000573	F-0VARC1000504	3903	R-0VARC1000504	8910
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	0VARC1000578 0VARC1000588	F-0VARC1000578	3905	R-0VARC1000578	8912
	0VARC1000588	F-0VARC1000588	3907	R-0VARC1000585	8913
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	0VARC1000649	F-0VARC1000649	3910	W-OAVWOIOOO40	0313
	0VARC1000649	F-0VARC1000661	3911	R-0VARC1000661	8916
40	0VARC1000678	F-0VARC1000678	3912	R-0VARC1000678	8917
	OVARC1000679	F-0VARC1000679	3913	R-nnnnnnnnnn	8918
	OVARC1000619	F-0VARC1000681	3914	R-0VARC1000681	8919
	0VARC1000682	F-0VARC1000682	3915	K OVARCIOOOOI	0313
45	0VARC1000682	F-0VARC1000689	3916	R-OVARC1000689	8920
	0VARC1000089	F-0VARC1000700	3917	R-OVARC1000700	8921
	0VARC1000700	F-0VARC1000700	3918	R-OVARC1000700	8922
	0VARC1000703	F-0VARC1000703	3919	K OTAKOTOOTOS	0322
50	0VARC1000722	F-0VARC1000722	3920	R-OVARC1000730	8923
	0VARC1000730	F-0VARC1000730	3920	R-OVARC1000730	8924
	0VARC1000748	F-0VARC1000748	3922	R-OVARC1000740	8925
	0VARC1000769 0VARC1000771	F-0VARC1000769 F-0VARC1000771	3922	R-0VARC1000709	8926
55	0VARC1000771	F-0VARC1000771	3923 3924	R-OVARC1000771	8927
-	0VARC1000781	F-0VARC1000787	3925	R-0VARC1000787	8928
	01WC1000191	I O'MICIUUUISI	JULU	W OAUWCTOOOLOL	0320

		EP 107	4011	AZ	
	OVARC1000800	F-0VARC1000800	3926	R-0VARC1000800	8929
	0VARC1000802	F-0VARC1000802	3927	R-0VARC1000802	8930
	0VARC1000834	F-0VARC1000834	3928	R-0VARC1000834	8931
5	0VARC1000846	F-0VARC1000846	3929	R-0VARC1000846	8932
	0VARC1000850	F-0VARC1000850	3930	R-0VARC1000850	8933
	0VARC1000862	F-0VARC1000862	3931	R-0VARC1000862	8934
	0VARC1000876	F-0VARC1000876	3932	R-0VARC1000876	8935
10	0VARC1000883	F-0VARC1000883	3933	R-0VARC1000883	8936
70	0VARC1000885	F-0VARC1000885	3934	R-0VARC1000885	8937
	0VARC1000886	F-0VARC1000886	3935	R-0VARC1000886	8938
	OVARC1000890	F-0VARC1000890	3936	K OMMOTOUGGO	0000
45	0VARC1000891	F-0VARC1000891	3937	R-0VARC1000891	8939
15	OVARC1000897	F-0VARC1000897	3938	R-OVARC1000897	8940
	OVARC1000831	F-OVARC1000897	3939	R-0VARC1000912	8941
	OVARC1000915	F-0VARC1000915	3940	R-0VARC1000915	8942
	OVARC1000913	F-OVARC1000913			
20	OVARC1000924 OVARC1000936		3941	R-0VARC1000924	8943
	OVARC1000936	F-OVARC1000936 F-OVARC1000937	3942	R-0VARC1000936	8944
	OVARC1000937	F-0VARC1000937	3943	R-0VARC1000937	8945
			3944	R-0VARC1000945	8946
25	OVARC1000948	F-0VARC1000948	3945	R-OVARC1000948	8947
	OVARC1000959 OVARC1000960	F-0VARC1000959	3946	R-OVARC1000959	8948
		F-0VARC1000960	3947	R-0VARC1000960	8949
	OVARC1000964	F-0VARC1000964	3948	D OUADOLOGO	0050
30	OVARC1000971	F-0VARC1000971	3949	R-0VARC1000971	8950
	OVARC1000984	F-0VARC1000984	3950	R-0VARC1000984	8951
	OVARC1000996	F-0VARC1000996	3951	R-0VARC1000996	8952
	OVARC1000999	F-0VARC1000999	3952	R-0VARC1000999	8953
35	OVARC1001000	F-0VARC1001000	3953	R-OVARC1001000	8954
30	OVARC1001004	F-0VARC1001004	3954	R-0VARC1001004	8955
	OVARC1001010	F-0VARC1001010	3955	R-0VARC1001010	8956
	0VARC1001011	F-0VARC1001011	3956	R-0VARC1001011	8957
	OVARC1001032	F-0VARC1001032	3957	R-0VARC1001032	8958
40	0VARC1001034	F-0VARC1001034	3958	R-0VARC1001034	8959
	OVARC1001038	F-0VARC1001038	3959	R-OVARC1001038	8960
	0VARC1001040	F-0VARC1001040	3960	R-OVARC1001040	8961
	0VARC1001044	F-0VARC1001044	3961	R-OVARC1001044	8962
45	OVARC1001051	F-0VARC1001051	3962	R-OVARC1001051	8963
	OVARC1001055	F-0VARC1001055	3963	R-0VARC1001055	8964
	OVARC1001062	F-0VARC1001062	3964	R-0VARC1001062	8965
	OVARC1001065	F-0VARC1001065	3965		
50	OVARC1001068	F-0VARC1001068	3966	R-0VARC1001068	8966
	OVARC1001072	F-0VARC1001072	3967	R-0VARC1001072	8967
	OVARC1001074	F-0VARC1001074	3968	R-0VARC1001074	8968
	OVARC1001085	F-0VARC1001085	3969	R-0VARC1001085	8969
55	OVARC1001092	F-0VARC1001092	3970	R-0VARC1001092	8970
	OVARC1001107	F-0VARC1001107	3971		

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		OVARC1001113	F-0VARC1001113	3972	R-0VARC1001113	8971
		OVARC1001117	F-0VARC1001117	3973	R-0VARC1001117	8972
		OVARC1001118	F-0VARC1001118	3974	R-0VARC1001118	8973
5		OVARC1001129	F-0VARC1001129	3975	R-0VARC1001129	8974
-		OVARC1001154	F-0VARC1001154	3976		
		OVARC1001161	F-0VARC1001161	3977	R-0VARC1001161	8975
		OVARC1001162	F-0VARC1001162	3978	R-0VARC1001162	8976
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10		OVARC1001169	F-0VARC1001169	3980	R-OVARC1001169	8978
		OVARC1001170	F-0VARC1001170	3981	R-0VARC1001170	8979
		OVARC1001171	F-0VARC1001171	3982	K OVIDIOTOOTTO	0010
		OVARC1001171	F-0VARC1001173	3983	R-0VARC1001173	8980
15		OVARC1001175	F-0VARC1001176	3984	K OVARCIOOII13	0300
		OVARC1001170	F-0VARC1001180	3985	R-0VARC1001180	8981
		OVARC1001180	F-0VARC1001180	3986	R-0VARC1001180	8982
20		OVARC1001200 OVARC1001232	F-0VARC1001200	3987	R-OVARC1001200	8983
			F-0VARC1001232	3988	R-OVARC1001232	8984
		OVARC1001240	F-0VARC1001240	3989	R-OVARC1001240	8985
		OVARC1001243	F-0VARC1001243	3990	R-OVARC1001243	8986
25		OVARC1001244	F-0VARC1001244	3991	D 01/4 D01 001 001	0007
		OVARC1001261	F-0VARC1001261	3992	R-0VARC1001261	8987
		OVARC1001268	F-0VARC1001268	3993	R-0VARC1001268	8988
		OVARC1001270	F-0VARC1001270	3994	R-0VARC1001270	8989
30		OVARC1001271	F-0VARC1001271	3995	R-OVARC1001271	8990
30		OVARC1001282	F-0VARC1001282	3996	R-0VARC1001282	8991
		OVARC1001296	F-0VARC1001296	3997	R-0VARC1001296	8992
	•	OVARC1001306	F-0VARC1001306	3998	R-nnnnnnnnnnnn	8993
		OVARC1001329	F-0VARC1001329	3999	R-OVARC1001329	8994
35		OVARC1001330	F-OVARC1001330	4000	R-OVARC1001330	8995
		0VARC1001339	F-OVARC1001339	4001	R-0VARC1001339	8996
		OVARC1001341	F-0VARC1001341	4002	R-0VARC1001341	8997
		OVARC1001342	F-OVARC1001342	4003	R-0VARC1001342	8998
40		0VARC1001344	F-0VARC1001344	4004	R-0VARC1001344	8999
		OVARC1001357	F-OVARC1001357	4005	R-0VARC1001357	9000
		OVARC1001360	F-0VARC1001360	4006	R-0VARC1001360	9001
		OVARC1001369	F-0VARC1001369	4007	R-OVARC1001369	9002
45		OVARC1001372	F-0VARC1001372	4008	R-0VARC1001372	9003
		OVARC1001376	F-0VARC1001376	4009	R-0VARC1001376	9004
		OVARC1001381	F-0VARC1001381	4010	R-0VARC1001381	9005
		OVARC1001391	F-0VARC1001391	4011	R-0VARC1001391	9006
50		0VARC1001399	F-OVARC1001399	4012	R-nnnnnnnnnnn	9007
00		OVARC1001417	F-OVARC1001417	4013	R-0VARC1001417	9008
		OVARC1001419	F-0VARC1001419	4014	R-OVARC1001419	9009
		OVARC1001425	F-0VARC1001425	4015	R-OVARC1001425	9010
		OVARC1001436	F-0VARC1001436	4016	R-OVARC1001436	9011
55		OVARC1001442	F-0VARC1001442	4017	R-OVARC1001442	9012

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	OVARC1001453	F-0VARC1001453	4018	R-OVARC1001453	- 9013
	OVARC1001476	F-0VARC1001476	4019	R-0VARC1001476	9014
	OVARC1001480	F-0VARC1001480	4020	R-0VARC1001480	9015
5	OVARC1001489	F-0VARC1001489	4021	R-0VARC1001489	9016
	OVARC1001496	F-0VARC1001496	4022	R-0VARC1001496	9017
	OVARC1001506	F-0VARC1001506	4023	R-0VARC1001506	9018
	OVARC1001525	F-0VARC1001525	4024	R-0VARC1001525	9019
10	OVARC1001542	F-0VARC1001542	4025	R-0VARC1001542	9020
, ,	0VARC1001547	F-0VARC1001547	4026	R-0VARC1001547	9021
	0VARC1001555	F-0VARC1001555	4027		
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15	OVARC1001600	F-0VARC1001600	4029	R-0VARC1001600	9023
13	OVARC1001610	F-0VARC1001610	4030	R-0VARC1001610	9024
	OVARC1001611	F-0VARC1001611,	4031	R-0VARC1001611	9025
	OVARC1001615	F-0VARC1001615	4032	R-0VARC1001615	9026
	OVARC1001668	F-0VARC1001668	4033	R-0VARC1001668	9027
20	OVARC1001702	F-0VARC1001702	4034	R-OVARC1001702	9028
	OVARC1001703	F-0VARC1001703	4035	R-OVARC1001703	9029
	OVARC1001711	F-0VARC1001711	4036	R-OVARC1001711	9030
	OVARC1001713	F-0VARC1001713	4037		
25	OVARC1001726	F-0VARC1001726	4038	R-0VARC1001726	9031
	OVARC1001731	F-0VARC1001731	4039	R-0VARC1001731	9032
	OVARC1001745	F-0VARC1001745	4040	R-0VARC1001745	9033
	OVARC1001762	F-0VARC1001762	4041	R-nnnnnnnnnnn	9034
30	OVARC1001766	F-0VARC1001766	4042	R-0VARC1001766	9035
	OVARC1001767	F-0VARC1001767	4043	R-nnnnnnnnnnn	9036
	OVARC1001768	F-0VARC1001768	4044	R-0VARC1001768	9037
	OVARC1001791	F-0VARC1001791	4045	R-OVARC1001791	9038
35	OVARC1001795	F-OVARC1001795	4046	R-OVARC1001795	9039
	OVARC1001802	F-0VARC1001802	4047	R-OVARC1001802	9040
	OVARC1001805	F-0VARC1001805	4048	R-OVARC1001805	9041
	OVARC1001809	F-0VARC1001809	4049		
40	OVARC1001812	F-0VARC1001812	4050	R-0VARC1001812	9042
	OVARC1001813	F-0VARC1001813	4051	R-0VARC1001813	9043
	OVARC1001820	F-0VARC1001820	4052	R-0VARC1001820	9044
	OVARC1001828	F-0VARC1001828	4053	R-0VARC1001828	9045
45	OVARC1001846	F-0VARC1001846	4054	R-0VARC1001846	9046
40	0VARC1001861	F-0VARC1001861	4055	R-OVARC1001861	9047
	OVARC1001873	F-0VARC1001873	4056	R-OVARC1001873	9048
	0VARC1001879	F-0VARC1001879	4057	R-0VARC1001879	9049
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50	OVARC1001883	F-0VARC1001883	4059	R-0VARC1001883	9051
	OVARC1001900	F-0VARC1001900	4060	R-0VARC1001900	9052
	OVARC1001901	F-0VARC1001901	4061	R-0VARC1001901	9053
	OVARC1001911	F-0VARC1001911	4062	R-0VARC1001911	9054
55	OVARC1001916	F-0VARC1001916	4063	R-0VARC1001916	9055

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	OVARC1001928	F-0VARC1001928	4064	R-0VARC1001928	9056
	OVARC1001942	F-0VARC1001942	4065	R-0VARC1001942	9057
	OVARC1001943	F-0VARC1001943	4066	R-0VARC1001943	9058
5	0VARC1001949	F-0VARC1001949	4067	R-0VARC1001949	9059
	0VARC1001950	F-0VARC1001950	4068	R-0VARC1001950	9060
	0VARC1001987	F-0VARC1001987	4069	R-0VARC1001987	9061
	OVARC1001989	F-0VARC1001989	4070	R-0VARC1001989	9062
10	0VARC1002044	F-0VARC1002044	4071	R-0VARC1002044	9063
	OVARC1002050	F-0VARC1002050	4072	R-0VARC1002050	9064
	OVARC1002066	F-0VARC1002066	4073	R-0VARC1002066	9065
	OVARC1002082	F-0VARC1002082	4074	R-OVARC1002082	9066
45	OVARC1002107	F-0VARC1002107	4075	R-OVARC1002107	9067
15	OVARC1002112	F-0VARC1002112	4076	1 0111101001101	•••
	OVARC1002112	F-0VARC1002127,	4077	R-0VARC1002127	9068
	0VARC1002128	F-0VARC1002138	4078	R-0VARC1002138	9069
	OVARC1002143	F-0VARC1002143	4079	R-0VARC1002143	9070
20	OVARC1002116	F-0VARC1002156	4080	R-0VARC1002156	9071
	OVARC1002158	F-0VARC1002158	4081	R-0VARC1002158	9072
	OVARC1002165	F-0VARC1002165	4082	R-0VARC1002165	9073
	OVARC1002182	F-0VARC1002182	4083	R-0VARC1002182	9074
25	PLACE1000004	F-PLACE1000004	4084	R-PLACE1000004	9075
	PLACE1000005	F-PLACE1000005	4085	R-PLACE1000005	9076
	PLACE1000007	F-PLACE1000007	4086	R-PLACE1000007	9077
	PLACE1000014	F-PLACE1000014	4087	R-PLACE1000014	9078
30	PLACE1000031	F-PLACE1000031	4088	R-PLACE1000031	9079
	PLACE1000040	F-PLACE1000040	4089	R-PLACE1000040	9080
	PLACE1000048	F-PLACE1000048	4090	R-PLACE1000048	9081
	PLACE1000050	F-PLACE1000050	4091	R-PLACE1000050	9082
35	PLACE1000061	F-PLACE1000061	4092	R-PLACE1000061	9083
	PLACE1000066	F-PLACE1000066	4093	R-PLACE1000066	9084
	PLACE1000078	F-PLACE1000078	4094	R-PLACE1000078	9085
	PLACE1000081	F-PLACE1000081	4095	R-PLACE1000081	9086
40	PLACE1000094	F-PLACE1000094	4096	R-PLACE1000094	9087
40	PLACE1000133	F-PLACE1000133	4097	R-PLACE1000133	9088
	PLACE1000142	F-PLACE1000142	4098	R-PLACE1000142	9089
	PLACE1000184	F-PLACE1000184	4099	R-PLACE1000184	9090
	PLACE1000185	F-PLACE1000185	4100	R-PLACE1000185	9091
45	PLACE1000213	F-PLACE1000213	4101	R-PLACE1000213	9092
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	PLACE1000308	F-PLACE1000308	4106		
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55	PLACE1000374	F-PLACE1000374	4109	R-PLACE1000374	9099

		EP 1 07	4 617 A	12	
	PLACE1000380	F-PLACE1000380	4110	R-PLACE1000380	9100 -
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	PLACE1000401	F-PLACE1000401	4112	R-PLACE1000401	9102
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	PLACE1000421	F-PLACE1000421	4115	R-PLACE1000421	9105
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	PLACE1000492	F-PLACE1000492	4121	R-PLACE1000492	9111
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	PLACE1000547	F-PLACE1000547	4123	R-PLACE1000547	9113
	PLACE1000547	F-PLACE1000562	4123	R-PLACE1000547	- 1
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55	PLACE1000979	F-PLACE1000979	4155	R-PLACE1000979	9144
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		EP IV	401/	42	
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	PLACE1001545	F-PLACE1001545	4202	R-PLACE1001545	9189
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	PLACE1001756	F-PLACE1001756	4225	R-PLACE1001756	9211
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	PLACE1001781	F-PLACE1001781	4228	R-PLACE1001781	9214
	PLACE1001701	F-PLACE1001799	4229	R-PLACE1001799	9215
0.5	PLACE1001810	F-PLACE1001810	4230	K TENCETOOT133	
35	PLACE1001817	F-PLACE1001817	4231	R-PLACE1001817	9216
	PLACE1001821	F-PLACE1001821	4232	R-PLACE1001821	9217
	PLACE1001844	F-PLACE1001844	4233	K 1 ENODIOCIOZI	<i>5</i> 21 .
	PLACE1001845	F-PLACE1001845	4234	R-PLACE1001845	9218
40	PLACE1001869	F-PLACE1001869	4235	R-PLACE1001869	9219
	PLACE1001897	F-PLACE1001897	4236	R-PLACE1001897	9220
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	PLACE1002004	F-PLACE1002004	4242	R 1 MODIO 1303	0000
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	PLACE1002052	F-PLACE1002052	4244	R-PLACE1002052	9227
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	PLACE1002072	F-PLACE1002072	4246	R-PLACE1002072	9229
55	PLACE1002073	F-PLACE1002073	4247	R-PLACE1002073	9230
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	PLACE1002157	F-PLACE1002157	4253	R-PLACE1002157	9236
	PLACE1002163	F-PLACE1002163	4254	R-PLACE1002163	9237
40	PLACE1002170	F-PLACE1002170	4255	K I MIODIOODIOO	3201
10	PLACE1002171	F-PLACE1002171	4256	R-PLACE1002171	9238
	PLACE1002111	F-PLACE1002205	4257	R-PLACE1002205	9239
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	PLACE1002768	F-PLACE1002768	4292	R-PLACE1002768	9273
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		F-PLACE1002775	4294		0075	
	PLACE1002782	F-PLACE1002782		M I Ditoblogation	9275	
	PLACE1002794	F-PLACE1002794		I(I THIOTH A CO.	9276	
_	PLACE1002811	F-PLACE1002811		If I mionion	9277	
5	PLACE1002815	F-PLACE1002815	4298	R-PLACE1002815	9278	
	PLACE1002816	F-PLACE1002816	4299	R-PLACE1002816	9279	
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10	PLACE1002851	F-PLACE1002851	4302	R-PLACE1002851	9282	
	PLACE1002853	F-PLACE1002853	4303	R-PLACE1002853	9283	ļ
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	PLACE100324			R-PLACE1003256	9308	
	PLACE100325				9309	
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	PLACE100333					
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	PLACE10033					
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	PLACE10033				9318	
55	PLACE10033	69 F-PLACE100336	9 4339	2 V-LFVCETO03302	. ,,,,,	
						

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5	PLACE1003394	F-PLACE1003394	4343		
3	PLACE1003401	F-PLACE1003401	4344	R-PLACE1003401	9322
	PLACE1003420	F-PLACE1003420	4345	R-PLACE1003420	9323
	PLACE1003454	F-PLACE1003454	4346	R-PLACE1003454	9324
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10	PLACE1003493	F-PLACE1003493	4348	R-PLACE1003493	9326
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	PLACE1005550	F-PLACE1005550	4530	R-PLACE1005550	9505
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	PLACE1005557	F-PLACE1005557	4532	R-PLACE1005557	9507
	PLACE1005574	F-PLACE1005574	4533	R-PLACE1005574	9508
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55	PLACE4000269	F-PLACE4000269	5212	R-PLACE4000269	10154
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40	Y79AA1001185	F-Y79AA1001185	5475	R-Y79AA1001185	10393
	Y79AA1001211	F-Y79AA1001211	5476	R-Y79AA1001211	10394
	Y79AA1001216	F-Y79AA1001216	5477	R-Y79AA1001216	10395
	Y79AA1001228	F-Y79AA1001228	5478	R-Y79AA1001228	10396
45	Y79AA1001233	F-Y79AA1001233	5479	R-Y79AA1001233	10397
	Y79AA1001236	F-Y79AA1001236	5480	R-Y79AA1001236	10398
	Y79AA1001281	F-Y79AA1001281	5481	R-Y79AA1001281	10399
	Y79AA1001299	F-Y79AA1001299	5482	R-Y79AA1001299	10400
50	Y79AA1001312	F-Y79AA1001312	5483	R-Y79AA1001312	10401
	Y79AA1001323	F-Y79AA1001323	5484	R-Y79AA1001323	10402
	Y79AA1001384	F-Y79AA1001384	5485	R-Y79AA1001384	10403
	Y79AA1001391	F-Y79AA1001391	5486	R-Y79AA1001391	10404
55	Y79AA1001394	F-Y79AA1001394	5487	R-Y79AA1001394	10405
	Y79AA1001402	F-Y79AA1001402	5488	R-Y79AA1001402	10406
	Y79AA1001493	F-Y79AA1001493	5489	R-Y79AA1001493	10407
				- 	

		LF 1	014 011	75	
	Y79AA1001511	F-Y79AA1001511	5490	R-Y79AA1001511	10408
	Y79AA1001533	F-Y79AA1001533	5491	R-Y79AA1001533	10409
	Y79AA1001541	F-Y79AA1001541	5492	R-nnnnnnnnnnn	10410
5	Y79AA1001548	F-Y79AA1001548	5493	R-Y79AA1001548	10411
	Y79AA1001555	F-Y79AA1001555	5494	R-Y79AA1001555	10412
	Y79AA1001581	F-Y79AA1001581	5495		
	Y79AA1001585	F-Y79AA1001585	5496	R-Y79AA1001585	10413
10	Y79AA1001594	F-Y79AA1001594	5497	R-Y79AA1001594	10414
	Y79AA1001603	F-Y79AA1001603	5498	R-Y79AA1001603	10415
	Y79AA1001613	F-Y79AA1001613	5499	R-Y79AA1001613	10416
	Y79AA1001647	F-Y79AA1001647	5500	R-Y79AA1001647	10417
15	Y79AA1001665	F-Y79AA1001665	5501	R-Y79AA1001665	10418
	Y79AA1001679	F-Y79AA1001679	5502	R-Y79AA1001679	10419
	Y79AA1001692	F-Y79AA1001692	5503	R-nnnnnnnnnnn	10420
	Y79AA1001696	F-Y79AA1001696	5504	R-Y79AA1001696	10421
20	Y79AA1001705	F-Y79AA1001705	5505	R-Y79AA1001705	10422
	Y79AA1001711	F-Y79AA1001711	5506	R-Y79AA1001711	10423
	Y79AA1001781	F-Y79AA1001781	5507	R-Y79AA1001781	10424
	Y79AA1001805	F-Y79AA1001805	5508	R-nnnnnnnnnnn	10425
25	Y79AA1001827	F-Y79AA1001827	5509	R-Y79AA1001827	10426
	Y79AA1001846	F-Y79AA1001846	5510	R-Y79AA1001846	10427
	Y79AA1001848	F-Y79AA1001848	5511	R-Y79AA1001848	10428
	Y79AA1001866	F-Y79AA1001866	5512	R-Y79AA1001866	10429
30 •	Y79AA1001874	F-Y79AA1001874	5513	R-Y79AA1001874	10430
	Y79AA1001875	F-Y79AA1001875	5514	R-Y79AA1001875	10431
	Y79AA1001923	F-Y79AA1001923	5515	R-Y79AA1001923	10432
	Y79AA1001963	F-Y79AA1001963	5516		
35	Y79AA1002027	F-Y79AA1002027	5517	R-Y79AA1002027	10433
	Y79AA1002083	F-Y79AA1002083	5518	R-Y79AA1002083	10434
	Y79AA1002089	F-Y79AA1002089	5519	R-Y79AA1002089	10435
	Y79AA1002093	F-Y79AA1002093	5520	R-Y79AA1002093	10436
40	Y79AA1002103	F-Y79AA1002103	5521	R-Y79AA1002103	10437
	Y79AA1002115	F-Y79AA1002115	5522	R-Y79AA1002115	10438
	Y79AA1002125	F-Y79AA1002125	5523	R-Y79AA1002125	10439
	Y79AA1002139	F-Y79AA1002139	5524	R-Y79AA1002139	10440
45	Y79AA1002204	F-Y79AA1002204	5525	R-Y79AA1002204	10441
	Y79AA1002208	F-Y79AA1002208	5526	R-nnnnnnnnnn	10442
	Y79AA1002209	F-Y79AA1002209	5527	R-Y79AA1002209	10443
	Y79AA1002210	F-Y79AA1002210	5528	R-Y79AA1002210	10444
50	Y79AA1002211	F-Y79AA1002211	5529	R-Y79AA1002211	10445
	Y79AA1002220	F-Y79AA1002220	5530	R-Y79AA1002220	10446
	Y79AA1002229	F-Y79AA1002229	5531	R-Y79AA1002229	10447
	Y79AA1002234	F-Y79AA1002234	5532	R-Y79AA1002234	10448
55	Y79AA1002246	F-Y79AA1002246	5533	R-Y79AA1002246	10449
,	Y79AA1002258	F-Y79AA1002258	5534	R-Y79AA1002258	10450
	Y79AA1002298	F-Y79AA1002298	5535	R-Y79AA1002298	10451

	Y79AA1002307	F-Y79AA1002307	5536	R-Y79AA1002307	10452	
	Y79AA1002311	F-Y79AA1002311	5537	R-Y79AA1002311	10453	
	Y79AA1002351	F-Y79AA1002351	5538	R-Y79AA1002351	10454	
5	Y79AA1002361	F-Y79AA1002361	5539	R-Y79AA1002361	10455	
	Y79AA1002399	F-Y79AA1002399	5540	R-Y79AA1002399	10456	
	Y79AA1002407	F-Y79AA1002407	5541	R-Y79AA1002407	10457	
10	Y79AA1002416	F-Y79AA1002416	5542	R-Y79AA1002416	10458	
	Y79AA1002431	F-Y79AA1002431	5543	R-Y79AA1002431	10459	
	Y79AA1002433	F-Y79AA1002433	5544	R-nnnnnnnnnnn	10460	
	Y79AA1002472	F-Y79AA1002472	5545	R-Y79AA1002472	10461	
15	Y79AA1002482	F-Y79AA1002482	5546	R-Y79AA1002482	10462	
	Y79AA1002487	F-Y79AA1002487	5547	R-Y79AA1002487	10463	

Table 2

	name of	name of	SEQ ID		SEQ ID
	clone	5'-end	of 5'-end	3'-end	of 3'-end
		sequence	sequence	sequence	sequence
	HEMBA1000497	F-HEMBA1000497	7 16111	R-HEMBA100049	7 16165
	HEMBA1001750	F-HEMBA1001750	16112	R-HEMBA100175	
	HEMBA1003854	F-HEMBA1003854	16113	R-HEMBA100385	4 16167
j	HEMBA1004193	F-HEMBA100419	3 16114	R-HEMBA100419	3 16168
	HEMBA1004860	F-HEMBA1004860	16115	R-HEMBA100486	0 16169
	HEMBA1005572	F-HEMBA100557	2 16116	R-HEMBA100557	2 16170
)	HEMBA1006038	F-HEMBA1006038	3 16117	R-HEMBA100603	8 16171
,	HEMBA1006092	F-HEMBA100609	2 16118	R-HEMBA100609	2 16172
	HEMBA1006406	F-HEMBA100640	6 16119	R-HEMBA100640	6 16173
	HEMBA1006650	F-HEMBA100665	0 16120	R-HEMBA100665	0 16174
5	HEMBA1006812	F-HEMBA100681	2 16121	R-HEMBA100681	2 16175
	HEMBB1000672	F-HEMBB100067	2 16122	R-HEMBB100067	2 16176
	HEMBB1001197	F-HEMBB100119	7 16123	R-HEMBB100119	7 16177
	HEMBB1001871	F-HEMBB100187	1 16124	R-HEMBB100187	71 16178
)	MAMMA1001252	F-MAMMA100125	2 16125	R-MAMMA100125	2 16179
	MAMMA1002094	F-MAMMA100209	4 16126	R-MAMMA100209	4 16180
	NT2RM4000634	F-NT2RM400063	4 16127	R-NT2RM400063	34 16181
_	NT2RM4000657	F-NT2RM400065	7 16128	R-NT2RM400069	57 16182
5	NT2RM4000783	F-NT2RM400078	3 16129	R-NT2RM400078	33 16183
	NT2RM4000857	F-NT2RM400085	7 16130	R-NT2RM40008	57 16184
	NT2RM4001178	F-NT2RM400117	8 16131	R-NT2RM40011	78 16185
)	NT2RM4002420	F-NT2RM400242	0 16132	R-NT2RM40024	20 16186
•	NT2RP2000198	F-NT2RP200019	8 16133	R-NT2RP20001	98 16187
	NT2RP2000551	F-NT2RP200055	1 16134	R-NT2RP20005	51 16188
	NT2RP2000660	F-NT2RP200066	0 16135	R-NT2RP20006	60 16189

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	NT2RP2001214	F-NT2RP2001214	16136	R-NT2RP2001214	16190
	NT2RP2001460	F-NT2RP2001460	16137	R-NT2RP2001460	16191
5	NT2RP2001756	F-NT2RP2001756	16138	R-NT2RP2001756	16192
5	NT2RP2002056	F-NT2RP2002056	16139	R-NT2RP2002056	16193
	NT2RP2002677	F-NT2RP2002677	16140	R-NT2RP2002677	16194
	NT2RP2002755	F-NT2RP2002755	16141	R-NT2RP2002755	16195
10	NT2RP2002843	F-NT2RP2002843	16142	R-NT2RP2002843	16196
	NT2RP2003101	F-NT2RP2003101	16143	R-NT2RP2003101	16197
	NT2RP2003799	F-NT2RP2003799	16144	R-NT2RP2003799	16198
	NT2RP2004095	F-NT2RP2004095	16145	R-NT2RP2004095	16199
15	NT2RP2004732	F-NT2RP2004732	16146	R-NT2RP2004732	16200
	NT2RP2004920	F-NT2RP2004920	16147	R-NT2RP2004920	16201
	NT2RP2005454	F-NT2RP2005454	16148	R-NT2RP2005454	16202
20	NT2RP2005776	F-NT2RP2005776	16149	R-NT2RP2005776	16203
20	NT2RP2005806	F-NT2RP2005806	16150	R-NT2RP2005806	16204
	NT2RP2005882	F-NT2RP2005882	16151	R-NT2RP2005882	16205
	NT2RP3001282	F-NT2RP3001282	16152	R-NT2RP3001282	16206
25	NT2RP3001723	F-NT2RP3001723	16153	R-NT2RP3001723	16207
	NT2RP3002099	F-NT2RP3002099	16154	R-NT2RP3002099	16208
	NT2RP3003155	F-NT2RP3003155	16155	R-NT2RP3003155	16209
	NT2RP3004028	F-NT2RP3004028	16156	R-NT2RP3004028	16210
30	0VARC1000008	F-0VARC1000008	16157	R-0VARC1000008	16211
	0VARC1000724	F-0VARC1000724	16158	R-0VARC1000724	16212
	OVARC1000751	F-0VARC1000751	16159	R-0VARC1000751	16213
35	OVARC1001029	F-0VARC1001029	16160	R-0VARC1001029	16214
00	PLACE1000814	F-PLACE1000814	16161	R-PLACE1000814	16215
	PLACE1003030	F-PLACE1003030	16162	R-PLACE1003030	16216
	PLACE1005549	F-PLACE1005549	16163	R-PLACE1005549	16217
40	PLACE1007218	F-PLACE1007218	16164	R-PLACE1007218	16218

Furthermore, the present invention relates to the use of the above primers, as described 45 below.

- (4) A polynucleotide which can be synthesized with the primer set of (2) or (3).
- 50 (5) A polynucleotide comprising a coding region in the polynucleotide of (4).
 - (6) A substantially pure protein encoded by polynucleotide of (4).
- 55 (7) A partial peptide of the protein of (6).

In addition, the present invention comprises a polynucleotide described below and a protein encoded by the polynucleotide.

5 (8) An isolated polynucleotide selected from the group consisting of

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- (a) a polynucleotide comprising a coding region of the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351;
- (b) a polynucleotide comprising a nucleotide sequence encoding a protein comprising the amino acid sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351;
- (c) a polynucleotide comprising a nucleotide sequence encoding a protein comprising an amino acid sequence selected from the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351, in which one or more amino acids are substituted, deleted, inserted, and/or added, wherein said protein is functionally equivalent to the protein comprising said amino acid sequence selected from the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351;
- (d) a polynucleotide that hybridizes with a polynucleotide comprising a nucleotide sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351, and that comprises a nucleotide sequence encoding a protein functionally equivalent to the protein encoded by the nucleotide sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351;
 - (e) a polynucleotide comprising a nucleotide sequence encoding a partial amino acid sequence of a protein encoded by the polynucleotide of (a) to (d);
- (f) a polynucleotide comprising a nucleotide sequence with at least 70% identity to the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351.
- 40 (9). A substantially pure protein encoded by the polynucleotide of (8).
 - (10) An antibody against the protein or peptide of any one of (6), (7), and (9).
- 45 (11) A vector comprising the polynucleotide of (5) or (8).
 - (12) A transformant carrying the polynucleotide of (5) or (8), or the vector of (11).
- (13) A transformant expressively carrying the polynucleotide of (5) or (8), or the vector of (11).
 - (14) A method for producing the protein or peptide of any one of (6), (7), and (9), comprising culturing the transformant of (13) and recovering the expression product.
 - (15) An oligonucleotide comprising the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351 or the nucleotide sequence complementary to the

complementary strand thereof, wherein said oligonucleotide comprises 15 nucleotides or more.

- 5 (16) Use of the oligonucleotide of (15) as a primer for synthesizing a polynucleotide.
 - (17) Use of the oligonucleotide of (15) as a probe for detecting a gene.
- 10 (18) An antisense polynucleotide against the polynucleotide of (8), or the portion thereof.
 - (19) A method for synthesizing a polynucleotide, the method comprising:
- a) synthesizing a complementary strand using a cDNA library as a template, and using the primer set of (2) or (3), or the primer of (16); and
 - b) recovering the synthesized product.

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- (20) The method of (19), wherein the cDNA library is obtainable by oligo-capping method.
- 25 (21) The method of (19), wherein the complementary strand is obtainable by PCR.
 - (22) A method for detecting the polynucleotide of (8), the method comprising:
- a) incubating a target polynucleotide with the oligonucleotide of (15) under the conditions where hybridization occurs, and
 - b) detecting the hybridization of the target polynucleotide with the oligonucleotide of (15).

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- (23) A database of polynucleotides and/or proteins, the database comprising information on at least one sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351 and/or the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351, or a medium on which the database is stored.
- Any patents, patent applications, and publications cited herein are incorporated by reference.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 shows the restriction maps of vectors pME18SFL3 and pUC19FL3.

Figure 2 shows the reproducibility of gene expression analysis. The respective intensities of gene expression observed in independent set of experiments are plotted in the vertical axis as well as in the horizontal axis.

Figure 3 shows the detection limit in gene expression analysis. The intensity of expression is shown in the vertical axis and the concentration (µg/ml) of probe used is shown in the horizontal axis.

DETAILED DESCRIPTION OF THE INVENTION

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Herein, "polynucleotide" is defined as a molecule in which multiple nucleotides are polymerized. There are no limitations in the number of the polymerized nucleotides. In case that the polymer contains relatively low number of nucleotides, it is also described as an "oligonucleotide". The polynucleotide or the oligonucleotide of the present invention can be a natural or chemically synthesized product. Alternatively, it can be synthesized using a template DNA by an enzymatic reaction such as PCR.

All the cDNA provided by the invention are full-length cDNA. Herein, a "full-length cDNA" is defined as a cDNA which contains both ATG codon (the translation start site) and the stop codon. Accordingly, the untranslated regions, which are originally found in the upstream or downstream of the protein coding region in natural mRNA, may or may not be contained.

An "isolated polynucleotide" is a polynucleotide the structure of which is not identical to that of any naturally occurring nucleic acid or to that of any fragment of a naturally occurring genomic nucleic acid spanning more than three separate genes. The term therefore covers, for example,

- (a) a DNA which has the sequence of part of a naturally occurring genomic DNA molecule but is not flanked by both of the coding sequences that flank that part of the molecule in the genome of the organism in which it naturally occurs;
- (b) a nucleic acid incorporated into a vector or into the genomic DNA of a prokaryote or eukaryote in a manner such that the resulting molecule is not identical to any naturally occurring vector or genomic DNA;
- (c) a separate molecule such as a cDNA, a genomic fragment, a fragment produced by polymerase chain reaction (PCR), or a restriction fragment; and
 - (d) a recombinant nucleotide sequence that is part of a hybrid gene, i.e., a gene encoding a fusion protein. Specifically excluded from this definition are nucleic acids present in mixtures of different (i) DNA molecules, (ii) transfected cells, or (iii) cell clones: e.g., as these occur in a DNA library such as a cDNA or genomic DNA library.

The term "substantially pure" as used herein in reference to a given polypeptide means that the protein or polypeptide is substantially free from other biological macromolecules. The substantially pure protein or polypeptide is at least 75% (e.g., at least 80, 85, 95, or 99%) pure by dry weight. Purity can be measured by any appropriate standard method, for example, by column chromatography, polyacrylamide gel electrophoresis, or HPLC analysis.

All the clones (5602 clones) of the present invention are novel and encode the full-length proteins. All the clones were prepared by oligo capping method, which can achieve cDNA cloning with high fullness ratio. The cDNA clones were selected by using ATGprl score as an index of the fullness ratio at the 5'-end, based on the sequence features of the 5'-end sequences. Selection was further carried out by searching GenBank database for EST sequences homologous to 5'-

end sequence of each clone by BLAST [S.F. Altschul, W. Gish, W. Miller, E.W. Myers & D.L. Lipman J. Mol. Biol., 215:403-410 (1990); W. Gish, & D.J. States, Nature Genet., 3:266-272 (1993)] and by considering the number of matching (identical) EST sequences or the number of continuous amino acids in the 5'-end sequence initiated from the initiation codon.

Moreover, the clones were turn out to be not identical to any of the known human mRNA (namely novel) by homology search using the 5'-end sequence.

The primers of the present invention, which are used for synthesizing full-length cDNA, are selected from the group comprising SEQ ID NO: 1-5547 (5'-primer), or SEQ ID NO: 5548-10463 (3'-primer). Further, the primers of the present invention, which are used for synthesizing full-length cDNA, are selected from SEQ ID NO: 16111-16164 (5'-primer), or SEQ ID NO: 16165-16218 (3'-primer). Some of the nucleotides include a known EST as its part. However, the primers of the present invention are novel in terms that the primers enable to synthesize full-length cDNA. Because the known ESTs lack important information on what part of cDNA the ESTs correspond to, it is impossible to design primers on the basis of the ESTs.

All the full-length cDNA of the present invention can be synthesized using a primer set comprising the nucleotide sequences selected from both the 5'-and 3'-end sequences, or a set comprising a primer based on the 5'-end sequence and an oligo-dT primer, by a method such as PCR (Current protocols in Molecular Biology (1987) Ausubel et al. edit, John Wiley & Sons, Section 6.1-6.4).

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Specifically, PCR can be performed using an oligonucleotide that has 15 nucleotides longer, and specifically hybridizes with the complementary strand of the polynucleotide that contains the nucleotide sequence selected from the 5'-end sequences shown in Table 1 and 2 (SEQ ID NO: 1-5547, or SEQ ID NO: 16111-16164), and an oligo-dT primer as a 5'-, and 3'-primer, respectively. The length of the primers is usually 15-100 bp, and favorably between 15-35 bp. In case of LA PCR, which is described below, the primer length of 25-35 bp may provide a good result.

A method to design a primer that enables a specific amplification based on the given nucleotide sequence is known to those skilled in the art (Current Protocols in Molecular Biology, Ausubel et al. edit, (1987) John Wiley & Sons, Section 6.1-6.4). In designing a primer based on the 5'-end sequence, the primer is designed so as that, in principle, the amplification products will include the translation start site. Accordingly, in case that a given 5'-end nucleotide sequence is the 5'- untranslated region (5'UTR), any part of the sequence can be used as a 5'-primer as far as the specificity toward the target cDNA is insured. The translation start site can be predicted using a known method such as the ATGpr as described below.

When synthesizing a polynucleotide, the target nucleotide sequence to be amplified can extend to several thousand bp in some cDNA. However, it is possible to amplify such a long nucleotides by using such as LA PCR (Long and Accurate PCR). It is advantageous to use LA PCR when synthesizing long DNA. In LA PCR, in which a special DNA polymerase having 3' →5' exonuclease activity is used, misincorporated nucleotides can be removed. Accordingly, accurate synthesis of the complementary strand can be achieved even with a long nucleotide sequence. By using LA PCR, it is reported that amplification of a nucleotide with 20 kb longer can be achieved under desirable condition (Takeshi Hayashi (1996) Jikken-Igaku Bessatsu, "Advanced Technologies in PCR" Youdo-sha).

A template DNA for synthesizing the cDNA of the present invention can be obtained by using cDNA libraries that are prepared by various methods. The full-length cDNA clones obtained here

are those with high fullness ratio, which were obtained using a combination of (1) a method to prepare a full-length-enriched cDNA library using the oligo-capping method, and (2) an estimation system for fullness using the 5'-end sequence (selection based on the estimation by the ATGpr after removing clones that are not-full-length compared to the ESTs). However, it is possible to easily obtain a full-length cDNA by using the primers that are provided by the present invention, not by the above described specialized method.

The problem with the cDNA libraries prepared by the known methods or commercially available is that mRNA contained in the libraries has very low fullness ratio. Thus, it is difficult to screen full-length cDNA clone directly from the library using ordinary cloning methods. The present invention has revealed a primer that is capable of synthesizing a full-length cDNA. If provided with primers, it is possible to synthesize a target full-length cDNA by using enzymatic reactions such as PCR. In particular, a full-length-enriched cDNA library, synthesized by methods such as oligo-capping, is desirable to synthesize a full-length cDNA with more reliability.

The 5'-end sequence of the full-length cDNA clones of the invention can be used to isolate the regulatory element of transcription including the promoter on the genome. By the spring of the year 2000, a rough draft of the human genome (analysis of human genomic sequence with lower accuracy), which covers 90% of the genome, is planned to be accomplished, and by the year 2003, analysis of the entire human genomic sequence is going to be finished. However, it is hard to analyze with software the transcription start sites on the human genome, in which long introns exist. By contrast, it is easy to specify the transcription start site on the genomic sequence using the 5'-end sequence of the full-length cDNA clone, thus it is easy to obtain the genomic region involved in transcription regulation, which includes the promoter that is contained in the upstream of the transcription start site.

The full-length cDNAs cloned in the present invention are classified into 13 groups, based on the data such as ATGpr1 score, by which the fullness ratio can be evaluated. Specifically, the 13 groups consist of; the below-mentioned groups (1)-(3), containing 3690 clones (Table 9), and the group (12), containing 3 clones, wherein ATGpr1 (score defined in the ATGpr program) is higher than 0.3; and the below-mentioned groups (4)-(11), containing 1857 clones (Table 10), and the group (13), containing 52 clones, wherein, although ATGpr1 is 0.3 or less, the clones are judged to be full-length from various viewpoints. Names of the clones belonging to the groups (1)-(13) are as indicated in Examples or below.

(1) 1516 clones

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Among the 3690 clones that have the maximal ATGpr1 score higher than 0.3, 1516 clones are novel full-length clones, in which at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

(2)377 clones

Among the 3690, 377 clones are novel full-length clones, in which the number of human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

(3) 1797 clones

Among the 3690, 1797 clones are novel full-length clones, in which the number of human EST having identical sequence at the 5'-end is not more than 20 (except the clones described above).

(4) 453 clones

Among the 1857 clones in which the maximal ATGpr1 score is 0.3 or less, the following 453 clones are estimated to be novel full-length clones since the clones have the maximal score 0.3 or more in the ATGpr2, and at least either of the sequences of their 5'- and 3'-ends, or both are not identical to those of any human EST. The ATGpr2 score is determined by using the ATGpr program with neglecting the information of the frequency of the six nucleotides contained within the sequence between the ATG codon and the stop codon (the maximal length is 300 nucleotides from the ATG codon) (Salamo A.A., Nishikawa T., and Swindells M.B. (1998) Bioinformatics, 14: 384-390; http://www.hri.co.jp/atgpr/). The ATGpr program for calculating the ATGpr2 score is described as the ATGpr2 program in the followings.

(5) 24 clones

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Among the 1857 clones, 24 clones are estimated to be full-length since their maximal ATGpr2 scores are higher than 0.3, and also novel, though they have low scores in ATGpr1 program, in which the number of the human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

(6) 65 clones

Among the 1857 clones, 65 clones are estimated to be full-length since, though they have low scores in both programs, ATGpr1 and ATGpr2, the scores are the maximum in comparison to those of the other clones in the same cluster (at least two clones). The clones are also novel, if at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

(7) 32 clones

Among the 1857 clones, 32 clones are estimated to be full-length since, though they have low scores in both programs, ATGpr1 and ATGpr2, the scores are the maximum in comparison to those of the other clones in the same cluster (at least two clones). The clones are also novel, if the number of the human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

(8) 36 clones

Among the 1857 clones, 36 clones are full-length, which were selected by assembling the sequences of the other clones or human EST, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

(9) 81 clones

Among the 1857 clones, 81 clones are full-length, which were selected by assembling the sequences of the other clones or human EST, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if the number of the human EST having identical sequence at the 5'-end is not more than 20 (other than the clones in which at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST).

(10) 938 clones

Among the 1857 clones, 938 clones are estimated to be full-length according to the fullness ratio shown in Table 4, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least the sequence of the 5'-end is not identical to those of

any human EST.

(11) 228 clones

Among the 1857 clones, 228 clones are estimated to be full-length according to the fullness ratio shown in Table 7, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least the sequence of the 3'-end is not identical to those of any human EST.

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(12) 3 clones

Three clones, HEMBA1006812, HEMBB1001871, and NT2RP3001282, whose maximal ATGpr1 values are higher than 0.3, are full-length and novel clones whose 5'-end sequences presumably contain a coding region which is initiated with ATG codon and which encodes 100 amino acids or more.

(13) 52 clones

20 The following 52 clones, which have maximal ATGpr1 values of 0.3 or less, are full-length with the fullness ratios shown in Table 4 although the fullness ratios are low: HEMBA1000497. HEMBA1001750, HEMBA1003854. HEMBA1004193. HEMBA1004860. HEMBA1005572, HEMBA1006038, HEMBA1006092, HEMBA1006406, HEMBA1006650, 25 HEMBB1000672. HEMBB1001197. MAMMA1001252. MAMMA1002094, NT2RM4000634, NT2RM4000657. NT2RM4000783. NT2RM4000857. NT2RM4001178, NT2RM4002420, NT2RP2000198. NT2RP2000551. NT2RP2000660. NT2RP2001214, NT2RP2001460, NT2RP2001756, NT2RP2002056, NT2RP2002677, NT2RP2002755, NT2RP2002843, NT2RP2003101, NT2RP2003799, NT2RP2004095, NT2RP2004732, NT2RP2004920, NT2RP2005454, NT2RP2005806, NT2RP2005882, NT2RP2005776, NT2RP3001723, NT2RP3002099, NT2RP3003155, NT2RP3004028, OVARC1000008, OVARC1000724, OVARC1000751. OVARC1001029. PLACE1000814, PLACE1003030, PLACE1005549, PLACE1007218, NT2RP4002298.

Moreover, the clones are novel clones whose 5'-end sequences presumably contain a coding region which is initiated with ATG codon and which encodes 50 amino acids or more. Among them, the following 20 clones is predicted to contain a coding region with 100 amino acids or more and should encode proteins:

HEMBA1000497, HEMBA1003854, HEMBA1004193, NT2RM4000657, NT2RM4001178, NT2RP2002677, NT2RP2001756, NT2RP2002755, NT2RP2002843, NT2RP2004095, NT2RP2004920, NT2RP2005806, NT2RP3002099, NT2RP3003155, OVARC1000724, OVARC1001029, PLACE1000814, PLACE1003030, PLACE1005549, PLACE1007218.

The protein encoded by the polynucleotide of the invention can be prepared as a recombinant protein or as a natural protein. For example, the recombinant protein can be prepared by inserting the polynucleotide encoding the protein of the invention into a vector, introducing the vector into an appropriate host cell and purifying the protein expressed within the transformed host cell, as described below. In contrast, the natural protein can be prepared, for example, by utilizing an affinity column to which an antibody against the protein of the invention (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 16.1-16.19) is attached. The antibody used for affinity purification may be either a

polyclonal antibody, or a monoclonal antibody. Alternatively, in vitro translation (See, for example, "On the fidelity of mRNA translation in the nuclease-treated rabbit reticulocyte lysate system." Dasso M.C., and Jackson R.J. (1989) Nucleic Acids Res. 17: 3129-3144) may be used for preparing the protein of the invention.

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Proteins functionally equivalent to the proteins of the present invention can be prepared based on the activities, which were clarified in the above-mentioned manner, of the proteins of the present invention. Using the biological activity possessed by the protein of the invention as an index, it is possible to verify whether or not a particular protein is functionally equivalent to the protein of the invention by examining whether or not the protein has said activity.

Proteins functionally equivalent to the proteins of the present invention can be prepared by those skilled in the art, for example, by using a method for introducing mutations into an amino acid sequence of a protein (for example, site-directed mutagenesis (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 8.1-8.5). Besides, such proteins can be generated by spontaneous mutations. The present invention comprises the proteins having one or more amino acids substitutions, deletions, insertions and/or additions in the amino acid sequences of the proteins of the present invention (Tables 350 and 351), as far as the proteins have the equivalent functions to those of the proteins identified in the present Examples described later.

There are no limitations in the number and sites of amino acid mutations, as far as the proteins maintain the functions thereof. The number of mutations is typically 30% or less, or 20% or less, or 10% or less, preferably within 5% or less, or 3% or less of the total amino acids, more preferably within 2% or less or 1 % or less of the total amino acids. From the viewpoint of maintaining the protein function, it is preferable that a substituted amino has a similar property to that of the original amino acid. For example, Ala, Val, Leu, Ile, Pro, Met, Phe and Trp are assumed to have similar properties to one another because they are all classified into a group of non-polar amino acids. Similarly, substitution can be performed among non-charged amino acid such as Gly, Ser, Thr, Cys, Tyr, Asn, and Gln, acidic amino acids such as Asp and Glu, and basic amino acids such as Lys, Arg, and His.

In addition, proteins functionally equivalent to the proteins of the present invention can be isolated by using techniques of hybridization or gene amplification known to those skilled in the art. Specifically, using the hybridization technique (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 6.3-6.4)), those skilled in the art can usually isolate a DNA highly homologous to the DNA encoding the protein identified in the present Example based on the identified nucleotide sequence (Tables 350 and 351) or a portion thereof and obtain the functionally equivalent protein from the isolated DNA. The present invention include proteins encoded by the DNAs hybridizing with the DNAs encoding the proteins identified in the present Example, as far as the proteins are functionally equivalent to the proteins identified in the present Example. Organisms from which the functionally equivalent proteins are isolated are illustrated by vertebrates such as human, mouse, rat, rabbit, pig and bovine, but are not limited to these animals.

Washing conditions of hybridization for the isolation of DNAs encoding the functionally equivalent proteins are usually "1 \times SSC, 0.1% SDS, 37°C"; more stringent conditions are "0.5 \times SSC, 0.1% SDS, 42°C"; and still more stringent conditions are "0.1 \times SSC, 0.1% SDS, 65°C". Alternatively, the following conditions can be given as hybridization conditions of the present invention. Namely, conditions in which the hybridization is done at "6 \times SSC, 40%

Formamide, 25°C", and the washing at "1 \times SSC, 55°C" can be given. More preferable conditions are those in which the hybridization is done at "6 \times SSC, 40% Formamide, 37°C", and the washing at "0.2 \times SSC, 55°C". Even more preferable are those in which the hybridization is done at "6 \times SSC, 50% Formamide, 37°C", and the washing at "0.1 \times SSC, 62°C". The more stringent the conditions of hybridization are, the more frequently the DNAs highly homologous to the probe sequence are isolated. Therefore, it is preferable to conduct hybridization under stringent conditions. Examples of stringent conditions in the present invention are, washing conditions of "0.5 \times SSC, 0.1% SDS, 42°C", or alternatively, hybridization conditions of "6 \times SSC, 40% Formamide, 37°C", and the washing at "0.2 \times SSC, 55°C". However, the above-mentioned combinations of SSC, SDS and temperature conditions are indicated just as examples. Those skilled in the art can select the hybridization conditions with similar stringency to those mentioned above by properly combining the above-mentioned or other factors (for example, probe concentration, probe length and duration of hybridization reaction) that determines the stringency of hybridization.

The amino acid sequences of proteins isolated by using the hybridization techniques usually exhibit high homology to those of the proteins of the present invention, which are shown in Tables 350 and 351. The present invention encompasses a polynucleotide comprising a nucleotide sequence that has a high identity to the nucleotide sequence of claim 8 (a).

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Furthermore, the present invention encompasses a peptide, or protein comprising an amino acid sequence that has a high identity to the amino acid sequence encoded by the polynucleotide of claim 8 (b). The term "high identity" indicates sequence identity of at least 40% or more;

preferably 60% or more; and more preferably 70% or more. Alternatively, more preferable is identity of 90% or more, or 93% or more, or 95% or more, furthermore, 97% or more, or 99% or more. The identity can be determined by using the BLAST search algorithm.

With the gene amplification technique (PCR) (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 6.3-6.4)) using primers designed based on the nucleotide sequence (Tables 350 and 351) or a portion thereof identified in the present Example, it is possible to isolate a DNA fragment highly homologous to the polynucleotide sequence or a portion thereof and to obtain functionally equivalent protein to a particular protein identified in the present Example based on the isolated DNA fragment.

The "percent identity" of two amino acid sequences or of two nucleic acids is determined using the algorithm of Karlin and Altschul (Proc. Natl. Acad. Sei. USA 87:2264-2268, 1990), modified as in Karlin and Altschul (Proc. Natl. Acad. Sei. USA 90:5873-5877, 1993). Such an algorithm is incorporated into the BLASTN and BLASTX programs of Altschul et al. (J. Mol. Biol.215:403-410, 1990). BLAST nucleotide searches are performed with the BLASTN program, score = 100, wordlength = 12. BLAST protein searches are performed with the BLASTX program, score = 50, wordlength = 3. When gaps exist between two sequences, Gapped BLAST is utilized as described in Altschul et al. (Nucleic Acids Res.25:3389-3402,1997). When utilizing BLAST and Gapped BLAST programs, the default parameters of the respective programs (e.g., BLASTX and BLASTN) are used. See http://www.ncbi.nlm.nih.gov.

The present invention also includes a partial peptide of the proteins of the invention. The partial peptide comprises a protein generated as a result that a signal peptide has been removed from a secretory protein. If the protein of the present invention has an activity as a receptor or a ligand, the partial peptide may function as a competitive inhibitor of the protein

and may bind to the receptor (or ligand). In addition, the present invention comprises an antigen peptide for raising antibodies. For the peptides to be specific for the protein of the invention, the peptides comprise at least 7 amino acids, preferably 8 amino acids or more, more preferably 9 amino acids or more, and even more preferably 10 amino acids or more. The peptide can be used for preparing antibodies against the protein of the invention, or competitive inhibitors of them, and also screening for a receptor that binds to the protein of the invention. The partial peptides of the invention can be produced, for example, by genetic engineering methods, known methods for synthesizing peptides, or digesting the protein of the invention with an appropriate peptidase.

The present invention also relates to a vector into which the DNA of the invention is inserted. The vector of the invention is not limited as long as it contains the inserted DNA stably. For example, if *E. coli* is used as a host, vectors such as pBluescript vector (Stratagene) are preferable as a cloning vector. To produce the protein of the invention, expression vectors are especially useful. Any expression vector can be used as far as it is capable of expressing the protein *in vitro*, in *E. coli*, in cultured cells, or *in vivo*. For example, pBEST vector (Promega) is preferable for *in vitro* expression, pET vector (Invitrogen) for *E. coli*, pME18S-FL3 vector (GenBank Accession No. AB009864) for cultured cells, and pME18S vector (Mol. Cell. Biol. (1988) 8: 466-472) for *in vivo* expression. To insert the DNA of the invention, ligation utilizing restriction sites can be performed according to the standard method (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.4-11.11).

The present invention also relates to a transformant carrying the vector of the invention. Any cell can be used as a host into which the vector of the invention is inserted, and various kinds of host cells can be used depending on the purposes. For strong expression of the protein in eukaryotic cells, COS cells or CHO cells can be used, for example.

Introduction of the vector into host cells can be performed, for example, by calcium phosphate precipitation method, electroporation method (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 9.1-9.9), lipofectamine method (GIBCO-BRL), or microinjection method, etc.

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The primer of the present invention can be used for synthesizing full-length cDNA, and also for the detection and/or diagnosis of the abnormality of the protein of the invention encoded by the full-length cDNA. For example, by utilizing polymerase chain reaction (genomic DNA-PCR, or RT-PCR) using the primer of the invention, DNA encoding the protein of the invention can be amplified. It is also possible to obtain the regulatory region of expression in the 5'-upstream by using PCR or hybridization since the transcription start site within the genomic sequence can be easily specified based on the 5'-end sequence of the full-length cDNA. The obtained genomic region can be used for detection and/or diagnosis of the abnormality of the sequence by RFLP analysis, SSCP, or direct sequencing.

Furthermore, the "polynucleotide having a length of at least 15 nucleotides, comprising a nucleotide sequence that is complementary to a polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs in Tables 350 and 351, or its complementary strand" includes an antisense polynucleotide for suppressing the expression of the protein of the invention. To exert the antisense effect, the antisense polynucleotide has a length of at least 15 bp or more, for example, 50 bp or more, preferably 100 bp or more, and more preferably 500 bp or more, and has a length of usually 3000 bp or less and preferably 2000 bp or less. The antisense DNA can be used in the gene therapy of the diseases that are caused by the

abnormality of the protein of the invention (abnormal function or abnormal expression). Said antisense DNA can be prepared, for example, by the phosphorothioate method ("Physicochemical properties of phosphorothioate oligodeoxynucleotides." Stein (1988) Nucleic Acids Res. 16: 3209-3221) based on the nucleotide sequence of the DNA encoding the protein (for example, the DNA set forth in any one of SEQ ID NOs in Tables 350 and 351).

The polynucleotide or antisense DNA of the present invention can be used in gene therapy, for example, by administrating it into a patient by the in vivo or ex vivo method with virus vectors such as retrovirus vectors, adenovirus vectors, and adeno-associated virus vectors, or non-virus vectors such as liposome.

The present invention also relates to antibodies that bind to the protein of the invention.

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There are no limitations in the form of the antibodies of the invention. They include polyclonal antibodies, monoclonal antibodies, or their portions that can bind to the protein of the invention. They also include antibodies of all classes. Furthermore, special antibodies such as humanized antibodies are also included.

The polyclonal antibody of the invention can be obtained according to the standard method by synthesizing an oligopeptide corresponding to the amino acid sequence and immunizing rabbits with the peptides (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.12-11.13). The monoclonal antibody of the invention can be obtained according to the standard method by purifying the protein expressed in *E. coli*, immunizing mice with the protein, and producing a hybridoma cell by fusing the spleen cells and myeloma cells (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.4-11.11).

The antibody binding to the protein of the present invention can be used for purification of the protein of the invention, and also for detection and/or diagnosis of the abnormalities of the expression and structure of the protein. Specifically, proteins can be extracted, for example, from tissues, blood, or cells, and the protein of the invention is detected by Western blotting, immunoprecipitation, or ELISA, etc. for the above purpose.

Furthermore, the antibody binding to the protein of the present invention can be utilized for treating the diseases that associates with the protein of the invention. If the antibodies are used for treating patients, human antibodies or humanized antibodies are preferable in terms of their low antigenicity. The human antibodies can be prepared by immunizing a mouse whose immune system is replaced with that of human ("Functional transplant of megabase human immunoglobulin loci recapitulates human antibody response in mice" Mendez M.J. et al. (1997) Nat. Genet. 15: 146-156). The humanized antibodies can be prepared by recombination of the hypervariable region of a monoclonal antibody (Methods in Enzymology (1991) 203: 99-121).

The cDNA of the present invention encodes the amino acid sequence of a protein which is predicted to have the function(s) described below based on the homology search of the GenBank and SwissProt. Specifically, for instance, as shown in EXAMPLES, searching a known gene or protein that is homologous to the partial sequence of the full-length cDNA of the invention (5602 clone) and referring the function of the gene and of the protein encoded by the gene make it possible to predict the function of the protein encoded by the cDNA of the invention. In this way, each of 1437 clones out of the 5602 full-length cDNA clones of the invention was predicted to encode a protein that was classified into one or more of the following categories.

Secretory or membrane protein (261 clones)

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Glycoprotein-associated protein (113 clones)

5 Signal transduction-associated protein (148 clones)

Transcription-associated protein (233 clones)

10 Disease-associated protein (437 clones)

Enzyme or metabolism-associated protein (301 clones)

15 Cell division- or cell proliferation-associated protein (74 clones)

Cytoskeleton-associated protein (92 clones)

20 RNA synthesis-associated protein (280 clones)

Nuclear protein (352 clones)

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25 Protein synthesis- or transport-associated protein (112 clones)

Cellular defense-associated protein (23 clones)

30 Development- or growth-associated protein (23 clones)

It is also possible to predict the protein function by looking into the amino acid sequence for the motifs such as the signal sequence, transmembrane region, nuclear translocation signal, glycosylation signal, phosphorylation site, Zinc finger motif, and SH3 domain. The programs, PSORT (Nakai K., and Kanehisa M. (1992) Genomics 14: 897-911), SOSUI (Hirokawa T. et al. (1998) Bioinformatics 14: 378-379) (Mitsui Information Developing Inc.), and MEMSAT (Jones D.T., Taylor W.R., and Thornton J.M. (1994) Biochemistry 33: 3038-3049) can be used to predict the existence of the signal sequence or transmembrane region. Alternatively, a partial amino acid sequence of the protein is fused with another protein such as GFP, the fusion protein is transfected into cultured cells, and the localization is analyzed to predict the function of the original protein.

Based on the determined nucleotide sequences of the full-length cDNAs obtained in the present invention, it is possible to predict more detailed functions of the proteins encoded by the cDNA clones, for example, by searching the databases such as GenBank, Swiss-Prot and UniGene for homologies of the cDNAs; or by searching the amino acid sequences deduced from the full-length cDNAs for signal sequences by using software programs such as PSORT, for transmembrane regions by using software programs such as SOSUI or for motifs by using software programs such as Pfam (http://www.sanger.ac.uk/Software/Pfam/index.shtm1) and PROSITE (http://www.expasy.ch/prosite/). As a matter of course, the functions are often predictable by using partial sequence information (preferably 300 nucleotides or more) instead of the full-length nucleotide sequences. However, the result of the prediction by using partial

nucleotide sequence does not always agree with the result obtained by using full-length nucleotide sequence, and thus, it is needless to say that the prediction of function is preferably performed based on the full-length nucleotide sequences.

- GenBank, Swiss-Prot and UniGene databases were searched for homologies of the full-length nucleotide sequences of the 4997 clones (see Example 18). The amino acid sequences deduced from the full-length nucleotide sequences were searched for functional domains by PSORT, SOSUI and Pfam. Prediction of functions of proteins encoded by the clones and the
- 10 categorization thereof were performed based on these results obtained.

HEMBA1006789,

HEMBA1006758,

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The following 798 clones were categorized into secretory and/or membrane proteins. HEMBA1000356, HEMBA1000518, HEMBA1000531, HEMBA1000637, HEMBA1000719,
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HEMBA1000817, HEMBA1000822, HEMBA1000852, HEMBA1000870, HEMBA1000991,

HEMBA1001052, HEMBA1001071, HEMBA1001085, HEMBA1001286, HEMBA1001351, HEMBA1001407, HEMBA1001446, HEMBA1001515, HEMBA1001557, HEMBA1001569,

HEMBA1001407, HEMBA1001446, HEMBA1001515, HEMBA1001557, HEMBA1001569, HEMBA1001661, HEMBA1001734, HEMBA1001746, HEMBA1001866, HEMBA1002125, HEMBA1002150, HEMBA1002166, HEMBA1002417, HEMBA1002462, HEMBA1002475,

HEMBA1002477, HEMBA1002486, HEMBA1002609, HEMBA1002659, HEMBA1002661, HEMBA1002780, HEMBA1002818, HEMBA1002876, HEMBA1002921, HEMBA1003071, HEMBA1003077, HEMBA1003079, HEMBA1003086, HEMBA1003096, HEMBA1003281, HEMBA1003286, HEMBA1003538, HEMBA1003711, HEMBA1003742, HEMBA1003803,

25 HEMBA1004055, HEMBA1004143, HEMBA1004146, HEMBA1004207, HEMBA1004341, HEMBA1004461, HEMBA1004577, HEMBA1004637, HEMBA1004752, HEMBA1004756, HEMBA1004850, HEMBA1004889, HEMBA1004923. HEMBA1004930. HEMBA1005029.

HEMBA1005035, HEMBA1005050, HEMBA1005552, HEMBA1005576, HEMBA1005581. HEMBA1005588, HEMBA1005616, HEMBA1005699. HEMBA1005991, HEMBA1006036, HEMBA1006038, HEMBA1006067, HEMBA1006173, HEMBA1006198, HEMBA1006293, HEMBA1006310, HEMBA1006492, HEMBA1006502, HEMBA1006583, HEMBA1006659,

HEMBA1006921,

HEMBA1006926,

35 HEMBA1006976, HEMBA1007203, HEMBA1007301, HEMBB1000037, HEMBB1000050, HEMBB1000054, HEMBB1000175, HEMBB1000317, HEMBB1000556, HEMBB1000593, HEMBB1000631, HEMBB1000763, HEMBB1000827, HEMBB1000915, HEMBB1000975,

HEMBB1001112, HEMBB1001151, HEMBB1001177, HEMBB1001302, HEMBB1001348, HEMBB1001564, HEMBB1001630, HEMBB1001871, HEMBB1001872, HEMBB1001925, HEMBB1001962, HEMBB1002042, HEMBB1002044. HEMBB1002142. HEMBB1002190. HEMBB1002193, HEMBB1002247, HEMBB1002383, HEMBB1002387, HEMBB1002550, HEMBB1002600, HEMBB1002692. MAMMA1000045. MAMMA1000129,

MAMMA1000133, MAMMA1000277, MAMMA1000278, MAMMA1000410, MAMMA1000416, MAMMA1000472, MAMMA1000672, MAMMA1000684, MAMMA1000714, MAMMA1000734, MAMMA1000778, MAMMA1000798, MAMMA1000842, MAMMA1000859, MAMMA1000897, MAMMA1000956,

MAMMA1001008, MAMMA1001030, MAMMA1001041, MAMMA1001073, MAMMA1001080, MAMMA1001139, MAMMA1001154, MAMMA1001322, MAMMA1001388, MAMMA1001411, MAMMA1001487. MAMMA1001751, MAMMA1001754, MAMMA1001771, MAMMA1002009, MAMMA1002427, MAMMA1002428, MAMMA1002461, MAMMA1002524, MAMMA1002573,

MAMMA1002598, MAMMA1002655, MAMMA1002684, MAMMA1002769, MAMMA1002844, MAMMA1002881, MAMMA1002890, MAMMA1002938, MAMMA1002947, MAMMA1003035, MAMMA1003089, MAMMA1003146, MAMMA1003150, NT2RM1000035, NT2RM1000037,

MAMMA1003089, MAMMA1003146, MAMMA1003150, NT2RM1000035, NT2RM1000037, NT2RM1000062, NT2RM1000080, NT2RM1000092, NT2RM1000131, NT2RM1000199,

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	NT2RM1000257,	NT2RM1000260,	NT2RM1000355,	NT2RM1000430,	NT2RM1000563,
	NT2RM1000648,	NT2RM1000742,	NT2RM1000770,	NT2RM1000800,	NT2RM1000811,
	NT2RM1000833,	NT2RM1000857,	NT2RM1000867,	NT2RM1000882,	NT2RM1000905,
5	NT2RM1001008,				
	NT2RM1001115,	NT2RM1001139,	NT2RM2000259,	NT2RM2000260,	NT2RM2000287,
	NT2RM2000395,	NT2RM2000402,	NT2RM2000407,	NT2RM2000422,	NT2RM2000490,
	NT2RM2000522,	NT2RM2000566,	NT2RM2000581,	NT2RM2000609,	NT2RM2000821,
10	NT2RM2001370,	NT2RM2001393,	NT2RM2001499,	NT2RM2001547,	NT2RM2001613,
	NT2RM2001648,	NT2RM2001659,	NT2RM2001671,	NT2RM2001688,	NT2RM2001698,
	NT2RM2001718,	NT2RM2001753,	NT2RM2001760,	NT2RM2001785,	NT2RM2001930,
	NT2RM2001950,	NT2RM2001997,	NT2RM2001998,	NT2RM2002049,	NT2RM2002145,
15	NT2RM4000233,	NT2RM4000433,	NT2RM4000457,	NT2RM4000486,	NT2RM4000496,
	NT2RM4000520,	NT2RM4000634,	NT2RM4000674,	NT2RM4000700,	NT2RM4000764,
	NT2RM4000778,	NT2RM4000795,	NT2RM4000820,	NT2RM4000857,	NT2RM4001032,
	NT2RM4001054,	NT2RM4001116,	NT2RM4001455,	NT2RM4001666,	NT2RM4001810,
20	NT2RM4001813,	NT2RM4001930,	NT2RM4001987,	NT2RM4002054,	NT2RM4002073,
	NT2RM4002145,	NT2RM4002146,	NT2RM4002189,	NT2RM4002194,	NT2RM4002251,
	NT2RM4002339,	NT2RM4002438,	NT2RM4002446,	NT2RM4002452,	NT2RM4002460,
	NT2RM4002493,	NT2RM4002558,	NT2RM4002565,	NT2RM4002571,	NT2RM4002594,
25	NT2RP1000130,	NT2RP1000191,	NT2RP1000326,	NT2RP1000358,	NT2RP1000413,
	NT2RP1000418,	NT2RP1000547,	NT2RP1000609,	NT2RP1000677,	NT2RP1000767,
	NT2RP1000782,	NT2RP1000856,	NT2RP1001113,	NT2RP1001247,	NT2RP1001286,
	NT2RP1001310,	NT2RP1001311,	NT2RP1001313,	NT2RP1001385,	NT2RP1001449,
30	NT2RP1001546,	NT2RP1001569,	NT2RP2000032,	NT2RP2000040,	NT2RP2000056,
	NT2RP2000070,	NT2RP2000091,	NT2RP2000114,	NT2RP2000120,	NT2RP2000173,
	NT2RP2000175,	NT2RP2000195,	NT2RP2000257,	NT2RP2000270,	NT2RP2000283,
25	NT2RP2000288,	NT2RP2000289,	NT2RP2000459,	NT2RP2000516,	NT2RP2000660,
35	NT2RP2000842,	NT2RP2000892,	NT2RP2001081,	NT2RP2001268,	NT2RP2001295,
	NT2RP2001366,	NT2RP2001378,	NT2RP2001576,	NT2RP2001581,	NT2RP2001597,
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	PLACE1000653,	PLACE1000755,	PLACE1001054,	PLACE1001062,	PLACE1001672,
35	PLACE1001692,	PLACE1001748,	PLACE1001781,	PLACE1001817,	PLACE1001869,
	PLACE1001989,	PLACE1002073,	PLACE1002598,	PLACE1002908,	PLACE1002991,
	PLACE1003174,	PLACE1003176,	PLACE1003709,	PLACE1003885,	PLACE1003888,
	PLACE1003903,	PLACE1003915,	PLACE1004270,	PLACE1004428,	PLACE1004437,
40	PLACE1004751,	PLACE1004804,	PLACE1004918,	PLACE1005243,	PLACE1005305,
	PLACE1005373,	PLACE1005656,	PLACE1005763,	PLACE1005804,	PLACE1005953,
	PLACE1005955,	PLACE1006011,	PLACE1006469,	PLACE1006534,	PLACE1006626,
	PLACE1006731,	PLACE1006819,	PLACE1006829,	PLACE1006878,	PLACE1007226,
45	PLACE1007416,	PLACE1007649,	PLACE1007706,	PLACE1007729,	PLACE1007954,
	PLACE1007958,	PLACE1008111,	PLACE1008275,	PLACE1008330,	PLACE1008643,
	PLACE1009094,	PLACE1009130,	PLACE1009444,	PLACE1009763,	PLACE1009861,
	PLACE1009992,	PLACE1009997,	PLACE1010096,	PLACE1010362,	PLACE1010481,
50	PLACE1010662,	PLACE1011046,	PLACE1011219,	PLACE1011229,	PLACE1011332,
	PLACE1011635,	PLACE1011923,	PLACE2000021,	PLACE2000034,	PLACE2000398,
	PLACE2000404,	PLACE2000438,	PLACE3000009,	PLACE3000020,	PLACE3000059,
	PLACE3000147,	PLACE3000339,	PLACE3000350,	PLACE4000063,	PLACE4000100,
55	PLACE4000401,	PLACE4000548,	PLACE4000654,	SKNMC1000050,	THYRO1000072,
	THYRO1000197,	THYRO1000288,	THYRO1000605,	THYRO1000662,	THYRO1000756,
	THYRO1000852,	THYRO1000926,	THYRO1000934,	THYRO1000951,	THYRO1000983,

			EP 10/401/ AZ		
	THYRO1001003,	THYRO1001287,	THYRO1001374,	THYRO1001406,	THYRO1001617,
	THYRO1001671,	THYRO1001738,	Y79AA1000782,	Y79AA1001048,	Y79AA1001233,
	Y79AA1001394,	Y79AA1001493,	Y79AA1001548,	Y79AA1001581,	Y79AA1001603,
5	Y79AA1001827,	Y79AA1002027,	Y79AA1002209,	Y79AA1002211,	Y79AA1002361,
	Y79AA1002416, H	HEMBA1005732, MAN	MMA1000402,		
	The following 2	217 clones presumabl	y belong to ATP- ar	nd/or GTP-binding p	roteins.
	HEMBA1000012,	HEMBA1000129,	HEMBA1000185,	HEMBA1000491,	HEMBA1000531,
10	HEMBA1001019,	HEMBA1001174,	HEMBA1001387,	HEMBA1001595,	HEMBA1001723,
	HEMBA1001913,	HEMBA1002161,	HEMBA1002212,	HEMBA1002876,	HEMBA1002997,
	HEMBA1003250,	HEMBA1003291,	HEMBA1003369,	HEMBA1003555	HEMBA1003560,
	HEMBA1004131,	HEMBA1004199,	HEMBA1004202,	HEMBA1004354,	HEMBA1004697,
15	HEMBA1005047,	HEMBA1005595,	HEMBA1007018,	HEMBA1007151,	HEMBB1000083,
	HEMBB1000226,	HEMBB1000264,	HEMBB1000632,	HEMBB1000725,	HEMBB1001294,
	HEMBB1002193,	MAMMA1000085,	MAMMA1000612,	MAMMA1000731,	MAMMA1000738,
	MAMMA1001038,		MAMMA1001768,	MAMMA1003127,	NT2RM1000187,
20	NT2RM1000388,	NT2RM1000702,	NT2RM1000772,	NT2RM1000924,	NT2RM2000469,
	NT2RM2000577,	NT2RM2000740,	NT2RM2001100,	NT2RM2001201,	NT2RM2001345,
	NT2RM2001823,	NT2RM2002128,	NT2RM4000155,	NT2RM4000191,	NT2RM4000356,
	NT2RM4000496	NT2RM4000611,	NT2RM4000733,	NT2RM4000820,	NT2RM4001084,
25	NT2RM4001178,	NT2RM4001344,	NT2RM4001444,	NT2RM4001592,	NT2RM4001714,
	NT2RM4001758,	NT2RM4001880,	NT2RM4002062,	NT2RM4002174,	NT2RM4002205,
	NT2RM4002527	NT2RM4002594,	NT2RM4002623,	NT2RP1000470,	NT2RP1000478.
	NT2RP1000915,	NT2RP1000958,	NT2RP1001080,	NT2RP1001410,	NT2RP1001569,
30	NT2RP2000126	NT2RP2000258,	NT2RP2000329,	NT2RP2000660,	NT2RP2000668,
	NT2RP2000710	NT2RP2000812,	NT2RP2000880,	NT2RP2001245,	NT2RP2001392,
	NT2RP2002606,	NT2RP2003277, NT		P2004538, NT2RP2	2004568,
	NT2RP2004689,	NT2RP2004768,	NT2RP2004791,	NT2RP2004920,	NT2RP2005344,
35	NT2RP2005393,	NT2RP2005763,	NT2RP2006534,	NT2RP3000046,	NT2RP3000252,
	NT2RP3000350,	NT2RP3000359,	NT2RP3000366,	NT2RP3000397,	NT2RP3000759,
	NT2RP3000845,	NT2RP3000875,	NT2RP3001150,	NT2RP3001427,	NT2RP3001453,
	NT2RP3001529,	NT2RP3001730,	NT2RP3001799,	NT2RP3001857,	NT2RP3001938,
40	NT2RP3002007,	NT2RP3002151,	NT2RP3002330,	NT2RP3002399,	NT2RP3002671,
	NT2RP3003301,	NT2RP3003353,	NT2RP3003589,	NT2RP3003809,	NT2RP3003876,
	NT2RP3004189,	NT2RP3004428,	NT2RP3004578,	NT2RP4000290,	NT2RP4000481,
	NT2RP4000518,	NT2RP4000781,	NT2RP4000839,	NT2RP4000929,	NT2RP4001041,
45	NT2RP4001079,	NT2RP4001375,	NT2RP4001414,	NT2RP4001592,	NT2RP4001634,
	NT2RP4001644,	NT2RP4001656,	NT2RP4001896,	NT2RP4002047,	NT2RP4002058,
	NT2RP4002408,	NT2RP5003477,	OVARC1000013,	OVARC1000304,	OVARC1000556,
	OVARC1000771,	OVARC1000800,	OVARC1001068,	OVARC1002138,	PLACE1000040,
50	PLACE1000588,	PLACE1001104,	PLACE1001739,	PLACE1002433,	PLACE1002437,
	PLACE1002714,	PLACE1003394,	PLACE1003521,	PLACE1003915,	PLACE1004902,
	PLACE1005243,	PLACE1005305,	PLACE1005549,	PLACE1005739,	PLACE1005921,
	PLACE1006119,	PLACE1006196,	PLACE1006552,	PLACE1006956,	PLACE1007409,
55	PLACE1007697,	PLACE1007946,	PLACE1008244,	PLACE1009404,	PLACE1009476,
	PLACE1009596,	PLACE1009908,	PLACE1010134,	PLACE1010720,	PLACE1010896,
	PLACE1011109,	PLACE1011114,	PLACE1011310,	PLACE1011922,	PLACE2000014,

	PLACE2000039,	PLACE2000274,	PLACE2000404,	PLACE2000427,	PLACE3000350,
	PLACE4000009,	PLACE4000014,	PLACE4000326,	SKNMC1000013,	THYRO1000072,
	THYRO1001458,	Y79AA1000833,	Y79AA1000962,	Y79AA1001394,	Y79AA1001875,
5	Y79AA1001963, Y	79AA1002209,			
	The following 3	20 clones presumab	ly belong to nuclear	proteins.	
	HEMBA1000005,	HEMBA1000158,	HEMBA1000216,	HEMBA1000561,	HEMBA1000591,
	HEMBA1001088,	HEMBA1001137,	HEMBA1001405,	HEMBA1001510,	HEMBA1001579,
10	HEMBA1001809,	HEMBA1001819,	HEMBA1001824,	HEMBA1001847,	HEMBA1001869,
	HEMBA1002177,	HEMBA1002241,	HEMBA1002495,	HEMBA1002569,	HEMBA1002935,
	HEMBA1002951,	HEMBA1002999,	HEMBA1003408,	HEMBA1003545,	HEMBA1003662,
	HEMBA1003684,	HEMBA1003690,	HEMBA1003760,	HEMBA1004203,	HEMBA1004321,
15	HEMBA1004353,	HEMBA1004479,	HEMBA1004973,	HEMBA1005219,	HEMBA1005359,
	HEMBA1005558,	HEMBA1005931,	HEMBA1006278,	HEMBA1006283,	HEMBA1006359,
	HEMBA1006485,	HEMBA1007087,	HEMBB1000226,	HEMBB1000789,	HEMBB1001011,
	HEMBB1001056,	HEMBB1001242,	HEMBB1001482,	HEMBB1001915,	HEMBB1002134,
20	HEMBB1002217,	MAMMA1000183,	MAMMA1000731,	MAMMA1001105,	MAMMA1001222,
	MAMMA1001260,	MAMMA1001633,	MAMMA1001743,	MAMMA1001837,	MAMMA1002617,
	MAMMA1002869,	MAMMA1002937,	MAMMA1003011,	NT2RM1000086,	NT2RM1000187,
	NT2RM1000666,	NT2RM1000885,	NT2RM1000894,	NT2RM1001059,	NT2RM1001092,
25	NT2RM2000013,	NT2RM2000588,	NT2RM2000624,	NT2RM2000735,	NT2RM2000740,
	NT2RM2001105,	NT2RM2001635,	NT2RM2001670,	NT2RM2001771,	NT2RM2001823,
	NT2RM2001936,	NT2RM2001989,	NT2RM2002004,	NT2RM2002088,	NT2RM2002091,
	NT2RM4000024,	NT2RM4000046,	NT2RM4000104,	NT2RM4000202,	NT2RM4000215,
30	NT2RM4000290,	NT2RM4000531,	NT2RM4000751,	NT2RM4000996,	NT2RM4001092,
	NT2RM4001140, I	NT2RM4001200, NT	2RM4001483, NT2R	M4001566, NT2RM	14001592,
	NITODIA ADDA COZ	NT2RM4001783,	NT2RM4001823,	NT2RM4001828,	NITODNAAOOAOEO
	NT2RM4001597,	11121(1114001700,	111211111111111111111111111111111111111	1112111114001020,	NT2RM4001858,
	NT2RM4001597, NT2RM4001979,	NT2RP1000035,	NT2RP1000111,	NT2RP1000493,	NT2RP1000574,
35					
35	NT2RM4001979, NT2RP1000630, NT2RP1001013,	NT2RP1000035,	NT2RP1000111,	NT2RP1000493, NT2RP1000958, NT2RP2000076,	NT2RP1000574,
35	NT2RM4001979, NT2RP1000630, NT2RP1001013, NT2RP2000153,	NT2RP1000035, NT2RP1000902,	NT2RP1000111, NT2RP1000915,	NT2RP1000493, NT2RP1000958,	NT2RP1000574, NT2RP1000966,
	NT2RM4001979, NT2RP1000630, NT2RP1001013, NT2RP2000153, NT2RP2000420,	NT2RP1000035, NT2RP1000902, NT2RP1001177,	NT2RP1000111, NT2RP1000915, NT2RP2000008,	NT2RP1000493, NT2RP1000958, NT2RP2000076, NT2RP2000258, NT2RP2001420,	NT2RP1000574, NT2RP1000966, NT2RP2000126, NT2RP2000297, NT2RP2001756,
35 40	NT2RM4001979, NT2RP1000630, NT2RP1001013, NT2RP2000153, NT2RP2000420, NT2RP2001869,	NT2RP1000035, NT2RP1000902, NT2RP1001177, NT2RP2000161, NT2RP2000931, NT2RP2002079,	NT2RP1000111, NT2RP1000915, NT2RP2000008, NT2RP2000248, NT2RP2001233, NT2RP2002270,	NT2RP1000493, NT2RP1000958, NT2RP2000076, NT2RP2000258, NT2RP2001420, NT2RP2002503,	NT2RP1000574, NT2RP1000966, NT2RP2000126, NT2RP2000297, NT2RP2001756, NT2RP2002591,
	NT2RM4001979, NT2RP1000630, NT2RP1001013, NT2RP2000153, NT2RP2000420, NT2RP2001869, NT2RP2002880,	NT2RP1000035, NT2RP1000902, NT2RP1001177, NT2RP2000161, NT2RP2000931, NT2RP2002079, NT2RP2002939,	NT2RP1000111, NT2RP1000915, NT2RP2000008, NT2RP2000248, NT2RP2001233, NT2RP2002270, NT2RP2002993,	NT2RP1000493, NT2RP1000958, NT2RP2000076, NT2RP2000258, NT2RP2001420, NT2RP2002503, NT2RP2003137,	NT2RP1000574, NT2RP1000966, NT2RP2000126, NT2RP2000297, NT2RP2001756, NT2RP2002591, NT2RP2003157,
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40	NT2RM4001979, NT2RP1000630, NT2RP1001013, NT2RP2000153, NT2RP2000420, NT2RP2001869, NT2RP2002880, NT2RP2003277, NT2RP2003912,	NT2RP1000035, NT2RP1000902, NT2RP1001177, NT2RP2000161, NT2RP2000931, NT2RP2002079, NT2RP2002939, NT2RP2003286, NT2RP2004013,	NT2RP1000111, NT2RP1000915, NT2RP2000008, NT2RP2000248, NT2RP2001233, NT2RP2002270, NT2RP2002993, NT2RP2003308, NT2RP2004187,	NT2RP1000493, NT2RP1000958, NT2RP2000076, NT2RP2000258, NT2RP2001420, NT2RP2002503, NT2RP2003137, NT2RP2003347, NT2RP2004689,	NT2RP1000574, NT2RP1000966, NT2RP2000126, NT2RP2000297, NT2RP2001756, NT2RP2002591, NT2RP2003157, NT2RP2003714, NT2RP2004920,
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40	NT2RM4001979, NT2RP1000630, NT2RP1001013, NT2RP2000153, NT2RP2000420, NT2RP2001869, NT2RP2002880, NT2RP2003277, NT2RP2003912, NT2RP2005393, NT2RP2005767,	NT2RP1000035, NT2RP1000902, NT2RP1001177, NT2RP2000161, NT2RP2002079, NT2RP2002939, NT2RP2003286, NT2RP20034013, NT2RP2005436, NT2RP2005776,	NT2RP1000111, NT2RP1000915, NT2RP2000008, NT2RP2000248, NT2RP2001233, NT2RP2002270, NT2RP2002993, NT2RP2003308, NT2RP2004187, NT2RP2005496, NT2RP2005933,	NT2RP1000493, NT2RP1000958, NT2RP2000076, NT2RP2000258, NT2RP2001420, NT2RP2003503, NT2RP2003137, NT2RP2003347, NT2RP2004689, NT2RP2005539, NT2RP2005542,	NT2RP1000574, NT2RP1000966, NT2RP2000126, NT2RP2000297, NT2RP2001756, NT2RP2003591, NT2RP2003714, NT2RP2003714, NT2RP2004920, NT2RP2005701, NT2RP2006043,
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40 45	NT2RM4001979, NT2RP1000630, NT2RP1001013, NT2RP2000153, NT2RP2000420, NT2RP2001869, NT2RP2003277, NT2RP2003277, NT2RP2003912, NT2RP2005393, NT2RP2005767, NT2RP2006436, NT2RP3000527,	NT2RP1000035, NT2RP1000902, NT2RP1001177, NT2RP2000161, NT2RP2002079, NT2RP2002939, NT2RP2003286, NT2RP2004013, NT2RP2005436, NT2RP2005436, NT2RP2005776, NT2RP3000031, NT2RP3000590,	NT2RP1000111, NT2RP1000915, NT2RP2000008, NT2RP2000248, NT2RP2001233, NT2RP2002270, NT2RP2002993, NT2RP2003308, NT2RP2004187, NT2RP2005496, NT2RP2005933, NT2RP2005933, NT2RP3000050, NT2RP3000603,	NT2RP1000493, NT2RP1000958, NT2RP2000076, NT2RP2000258, NT2RP2001420, NT2RP2003137, NT2RP2003347, NT2RP2004689, NT2RP2005539, NT2RP2005942, NT2RP3000397, NT2RP3000632,	NT2RP1000574, NT2RP1000966, NT2RP2000126, NT2RP2000297, NT2RP2001756, NT2RP2003157, NT2RP2003714, NT2RP2004920, NT2RP2005701, NT2RP2006043, NT2RP3000512, NT2RP3000917,
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	NT2RP3002151,	NT2RP3002529,	NT2RP3002671,	NT2RP3003301,	NT2RP3003846,
	NT2RP3003876,	NT2RP3004209,	NT2RP4000370,	NT2RP4000457,	NT2RP4000879,
	NT2RP4000927,	NT2RP4001041,	NT2RP4001117,	NT2RP4001313,	NT2RP4001315,
45	NT2RP4001574,	NT2RP4001592,	OVARC1000013,	OVARC1000071,	OVARC1000085,
	OVARC1000465,	OVARC10005	64, OVARC10	00771, OVAF	RC1000862,
	OVARC1001171,	OVARC1001180,	OVARC1001342,	PLACE1000007,	PLACE1000061,
	PLACE1000081,	PLACE1000492,	PLACE1000863,	PLACE1001092,	PLACE1001748,
50	PLACE1002090,	PLACE1003174,	PLACE1003915,	PLACE1004104,	PLACE1004270,
	PLACE1004743,	PLACE1005557,	PLACE1005813,	PLACE1006170,	PLACE1006488,
	PLACE1006829,	PLACE1007706,	PLACE1007729,	PLACE1008273,	PLACE1008402,
	PLACE1008790,	PLACE1008813,	PLACE1009094	PLACE1009130,	PLACE1009477,
55	PLACE1009721,	PLACE1009845,	PLACE1010074,	PLACE1010547,	PLACE1011109,
	PLACE1011229,	PLACE1011477,	PLACE1012031,	PLACE2000404,	PLACE3000059,
	PLACE3000121,	PLACE4000269,	PLACE4000654,	SKNMC1000011,	THYRO1000983,
		•	·	•	·

THYRO1001003, THYRO1001313, Y79AA1000560, Y79AA1000784, Y79AA1000968, Y79AA1001493, Y79AA1001875, Y79AA1002027, Y79AA1002209, HEMBA1006284,

The following 130 clones presumably belong to cytoskeleton-associated proteins.

```
5
    HEMBA1000156,
                    HEMBA1000168,
                                     HEMBA1000411,
                                                     HEMBA1000588,
                                                                     HEMBA1001043.
    HEMBA1001651.
                     HEMBA1001661.
                                     HEMBA1002102,
                                                     HEMBA1002161,
                                                                     HEMBA1002939,
    HEMBA1003235,
                     HEMBA1003581,
                                     HEMBA1004499,
                                                     HEMBA1004534,
                                                                      HEMBA1004697,
    HEMBA1004929,
                     HEMBA1004972,
                                     HEMBA1005582,
                                                     HEMBA1005595,
                                                                      HEMBA1006344,
    HEMBA1006737,
                     HEMBB1001175,
                                     HEMBB1001282,
                                                     HEMBB1001562,
                                                                      HEMBB1001802,
    MAMMA1000824,
                    MAMMA1001041,
                                     MAMMA1001576,
                                                     MAMMA1001679,
                                                                     MAMMA1001735,
    MAMMA1002297,
                    MAMMA1002351,
                                     MAMMA1002622.
                                                     MAMMA1002637.
                                                                     MAMMA1003127.
    NT2RM1000850,
                     NT2RM1000898,
                                     NT2RM2000030,
                                                     NT2RM2000260,
                                                                      NT2RM2000691,
15
    NT2RM2001324,
                     NT2RM4000169.
                                     NT2RM4000229,
                                                     NT2RM4000515,
                                                                      NT2RM4001217,
    NT2RP1000202, NT2RP1000348, NT2RP1000460,
    NT2RP1000478,
                     NT2RP1001033,
                                     NT2RP1001294,
                                                      NT2RP1001302,
                                                                      NT2RP2000070,
    NT2RP2000812,
                     NT2RP2000814,
                                     NT2RP2001168,
                                                      NT2RP2001245,
                                                                      NT2RP2001634,
20
    NT2RP2001900,
                     NT2RP2003307,
                                     NT2RP2003394,
                                                      NT2RP2004041,
                                                                      NT2RP2004242,
    NT2RP2004538,
                     NT2RP2004587,
                                     NT2RP2004681,
                                                      NT2RP2004732,
                                                                      NT2RP2004978,
    NT2RP2005491,
                     NT2RP2005531,
                                     NT2RP2005712,
                                                      NT2RP2006275,
                                                                      NT2RP3000753,
    NT2RP3001113,
                     NT2RP3001216,
                                     NT2RP3001239,
                                                      NT2RP3001272,
                                                                      NT2RP3001554,
    NT2RP3001690,
                     NT2RP3001799,
                                     NT2RP3002688,
                                                      NT2RP3003061,
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    NT2RP3003230.
                     NT2RP3004569,
                                     NT2RP3004578,
                                                     NT2RP4001004,
                                                                      NT2RP4001086.
    NT2RP4001256,
                    NT2RP4001567,
                                     NT2RP4001927,
                                                     OVARC1000001,
                                                                      OVARC1000106,
                     OVARC1000520,
                                     OVARC1000679,
    OVARC1000437,
                                                     OVARC1001731,
                                                                      OVARC1002050,
    PLACE1001104, PLACE1002571,
    PLACE1002591,
                     PLACE1002655,
                                     PLACE1002714,
                                                     PLACE1003625,
                                                                      PLACE1005287,
    PLACE1006552,
                     PLACE1007946,
                                     PLACE1008426,
                                                      PLACE1010148,
                                                                      PLACE1010547,
    PLACE1010743,
                     PLACE1010896,
                                     PLACE1010960,
                                                      PLACE1011310,
                                                                      PLACE1011922,
    PLACE2000216,
                     PLACE2000274,
                                     PLACE2000371,
                                                      PLACE2000458,
                                                                      PLACE3000145,
    PLACE3000416,
                     PLACE4000009,
                                     THYRO1000132,
                                                     THYRO1001405,
                                                                      THYRO1001458,
    Y79AA1000368, Y79AA1000794, Y79AA1000833, Y79AA1000962, Y79AA1002208,
```

The following 54 clones presumably belong to cell division-associated and/or cell proliferation-associated proteins.

HEMBA1001019, HEMBA1001595, HEMBA1002363, HEMBA1002997, HEMBA1003136, HEMBA1003369, HEMBA1004131, HEMBA1004354, HEMBA1005621, HEMBB1000037, HEMBB1000264, MAMMA1001768, MAMMA1002769, NT2RM1000354, NT2RM1000430, NT2RM1000874, NT2RM2001256, NT2RM2001743, NT2RM2001896, NT2RM2002145, NT2RM4000215. NT2RM4001714. NT2RP1000163. NT2RP1000333, NT2RP1000439, NT2RP2000346, NT2RP2001397, NT2RP2002595, NT2RP2003177, NT2RP2003596 NT2RP2003912, NT2RP2004396, NT2RP2005037. NT2RP2005520, NT2RP2005669, NT2RP2005835, NT2RP3001730, NT2RP3002081, NT2RP4000210 NT2RP4000415, NT2RP4001414, NT2RP4001634, OVARC1000013, OVARC1000937, PLACE1001383, PLACE1002433, PLACE1004316, PLACE1005287. PLACE1008808. PLACE1010720, PLACE1010833, Y79AA1000748, Y79AA1001236, Y79AA1001394

The following 36 clones presumably belong to the category of embryogenesis- and/or development-associated proteins:

```
HEMBA1000518,
                     HEMBA1001847.
                                     HEMBA1001869,
                                                      HEMBA1003545,
                                                                      HEMBA1004973,
    HEMBB1002442,
                     MAMMA1001837,
                                      NT2RM2001670,
                                                      NT2RM4000046.
                                                                       NT2RM4000531,
                     NT2RM4001858,
                                      NT2RP2002078,
                                                      NT2RP2004187
    NT2RM4001140,
                                                                       NT2RP2006436,
    NT2RP3000603,
                     NT2RP3000994,
                                      NT2RP3001580,
                                                      NT2RP3001708.
                                                                       NT2RP3003071.
    NT2RP3004472,
                     NT2RP3004617,
                                     NT2RP4000246,
                                                      NT2RP4001567,
                                                                       OVARC1000304,
                                      PLACE1002532,
    OVARC1000746,
                     PLACE1000793,
                                                      PLACE1003258,
                                                                       PLACE1003625,
    PLACE1004460,
                     PLACE1009622,
                                      PLACE4000558,
                                                      THYRO1000085,
                                                                       Y79AA1001391,
10
    Y79AA1001692
                                     belong to cellular
      The following 30 clones presumably
                                                     defense-associated proteins.
    HEMBA1000005,
                     HEMBA1000531,
                                     HEMBA1003417,
                                                      HEMBA1006253.
                                                                       NT2RM4000354.
                     NT2RP1000333,
                                      NT2RP1000493,
                                                      NT2RP2000006.
    NT2RM4001880.
                                                                       NT2RP2000045,
15
                                      NT2RP2002464.
    NT2RP2000809,
                     NT2RP2001536,
                                                      NT2RP2004920,
                                                                       NT2RP2005037,
    NT2RP3000590,
                     NT2RP3001426,
                                      NT2RP3002062,
                                                      NT2RP3002785
                                                                       NT2RP3004262,
    NT2RP4001555,
                     NT2RP4001638,
                                      PLACE1006958,
                                                      PLACE1008275
                                                                       PLACE1009113,
    PLACE1011858, PLACE4000014, THYRO1000684, Y79AA1002139, Y79AA1002229
20
      Although it is unclear whether or not 261 clones out of clones other than the above-mentioned
    clones belong to any of the above-described categories, these clones are predicted to have
    some functions, based on the homology search using their full-length sequences
                     HEMBA1000307,
                                     HEMBA1000333,
                                                      HEMBA1000488,
    HEMBA1000030,
                                                                      HEMBA1000523,
    HEMBA1001197,
                     HEMBA1001302,
                                     HEMBA1001455,
                                                      HEMBA1001675,
                                                                      HEMBA1001714,
    HEMBA1001744.
                     HEMBA1001967.
                                     HEMBA1002151.
                                                      HEMBA1002215.
                                                                       HEMBA1002458.
    HEMBA1002777,
                     HEMBA1003098,
                                     HEMBA1003199,
                                                      HEMBA1003615,
                                                                      HEMBA1003836
    HEMBA1004295,
                     HEMBA1004573,
                                     HEMBA1004604,
                                                      HEMBA1004795,
                                                                       HEMBA1005101,
    HEMBA1005201,
                     HEMBA1005206
                                     HEMBA1005530,
                                                      HEMBA1005666,
                                                                       HEMBA1005990,
                     HEMBA1006398,
                                     HEMBA1006445,
                                                      HEMBA1007174,
                                                                       HEMBA1007251,
    HEMBA1006268,
    HEMBB1000036,
                     HEMBB1000144.
                                     HEMBB1000973.
                                                      HEMBB1001058.
                                                                       HEMBB1001234.
                     HEMBB1001331,
                                     HEMBB1001384,
                                                      HEMBB1002266,
                                                                       HEMBB1002510,
    HEMBB1001288,
    HEMBB1002705,
                    MAMMA1000055,
                                     MAMMA1000625,
                                                      MAMMA1001075,
                                                                      MAMMA1001181,
                                     MAMMA1002143,
    MAMMA1001259,
                     MAMMA1001730,
                                                      MAMMA1002699,
                                                                      MAMMA1002972,
                     NT2RM1000118,
                                      NT2RM1000186,
    MAMMA1003113,
                                                      NT2RM1000244,
                                                                       NT2RM1000421,
    NT2RM1000499,
                     NT2RM1000623,
                                      NT2RM1000883,
                                                      NT2RM2000502
                                                                       NT2RM2000599,
    NT2RM2000718,
                     NT2RM2001065,
                                      NT2RM2001196.
                                                      NT2RM2001983.
                                                                       NT2RM2002109.
    NT2RM2002142,
                     NT2RM4000030,
                                      NT2RM4000139,
                                                      NT2RM4000156,
                                                                       NT2RM4000386,
    NT2RM4000590,
                     NT2RM4001047,
                                      NT2RM4001155.
                                                      NT2RM4001256,
                                                                       NT2RM4001320,
    NT2RM4001340,
                     NT2RM4001347,
                                      NT2RM4001371,
                                                      NT2RM4001582,
                                                                       NT2RM4001611,
    NT2RM4001731,
                     NT2RM4001969,
                                      NT2RM4002034,
                                                      NT2RM4002075,
                                                                       NT2RM4002226,
    NT2RP1000040,
                     NT2RP1000363,
                                      NT2RP1000481.
                                                      NT2RP1000513,
                                                                       NT2RP1000733.
    NT2RP1000860, NT2RP1000954, NT2RP1001011, NT2RP1001395, NT2RP1001457,
    NT2RP1001494,
                     NT2RP2000054,
                                      NT2RP2000067,
                                                      NT2RP2000133,
                                                                       NT2RP2000157,
    NT2RP2000764,
                     NT2RP2000965.
                                      NT2RP2001839,
                                                      NT2RP2001883,
                                                                       NT2RP2001976,
    NT2RP2001985,
                     NT2RP2002185,
                                      NT2RP2002442,
                                                      NT2RP2002727,
                                                                       NT2RP2002741,
                                      NT2RP2003265,
                                                                       NT2RP2003857,
    NT2RP2002986,
                                                      NT2RP2003272,
                     NT2RP2003121,
    NT2RP2003871,
                     NT2RP2004425,
                                      NT2RP2004476,
                                                      NT2RP2004710,
                                                                       NT2RP2004816,
    NT2RP2005441,
                        NT2RP2005490,
                                            NT2RP2005620,
                                                                NT2RP2005654.
    NT2RP2005675,
                     NT2RP2005753,
                                      NT2RP2005841,
                                                      NT2RP2006598,
                                                                       NT2RP3000047,
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	NT2RP3000233,	NT2RP3000868,	NT2RP3000869,	NT2RP3001399,	NT2RP3001407,
	NT2RP3001457,	NT2RP3001587,	NT2RP3001712,	NT2RP3001819,	NT2RP3001854,
	NT2RP3001931,	NT2RP3002273,	NT2RP3002631,	NT2RP3002682,	NT2RP3002770,
5	NT2RP3002818,	NT2RP3002948,	NT2RP3002972,	NT2RP3003032,	NT2RP3003290,
	NT2RP3003411,	NT2RP3003491,	NT2RP3003500,	NT2RP3003726,	NT2RP3004348,
	NT2RP3004507,	NT2RP4000129,	NT2RP4000498,	NT2RP4000528,	NT2RP4000737,
	NT2RP4000979,	NT2RP4001010,	NT2RP4001207,	NT2RP4001228,	NT2RP4001260,
10	NT2RP4001339,	NT2RP4001351,	NT2RP4001474,	NT2RP4001966,	NT2RP4002018,
	OVARC1000209,	OVARC1000876,	OVARC1001065,	OVARC1001092,	OVARC1001419,
	OVARC1001555,	OVARC1001711,	OVARC1001943,	PLACE1000004,	PLACE1000066,
	PLACE1000610,	PLACE1000636,	PLACE1000769,	PLACE1000987,	PLACE1001036,
15	PLACE1001845,	PLACE1001920,	PLACE1002665,	PLACE1003602,	PLACE1003611,
	PLACE1004256,	PLACE1004550,	PLACE1004868,	PLACE1004930,	PLACE1005052,
	PLACE1005102,	PLACE1005176,	PLACE1005187,	PLACE1005331,	PLACE1005727,
	PLACE1006003,	PLACE1006335,	PLACE1006385,	PLACE1006506,	PLACE1007105,
20	PLACE1007537,	PLACE1007705,	PLACE1007791,	PLACE1007897,	PLACE1008080,
	PLACE1008368,	PLACE1008398,	PLACE1008465,	PLACE1008627,	PLACE1009020,
	PLACE1009060,	PLACE1009186,	PLACE1009443,	PLACE1009571,	PLACE1009670,
	PLACE1010105,	PLACE1010261,	PLACE1010310,	PLACE1010522,	PLACE1010579,
25	PLACE1010628,	PLACE1010661,	PLACE1010761,	PLACE1011185,	PLACE1011340,
	PLACE1011586,	PLACE2000246,	PLACE2000411,	PLACE3000477,	THYRO1000173,
	THYRO1000401,	THYRO1000666,	THYRO1001033,	THYRO1001347,	THYRO1001656,
	THYRO1001703,	THYRO1001721,	Y79AA1000059,	Y79AA1000181,	Y79AA1000268,
30	Y79AA1000313,	Y79AA1000540,	Y79AA1000966,	Y79AA1000985,	Y79AA1001323,
	Y79AA1001402,	Y79AA1001679,	Y79AA1001923,	Y79AA1002083,	Y79AA1002307,
	Y79AA1002311, Y	79AA1002487,			

In some cases, the predicted functions based on the partial sequences are different from those based on the full-length sequences. The reason is that a protein does not always belong solely to a single category of the above-described functional categories, and therefore, a protein may belong to two or more of the predicted functional categories. Besides, additional functions can be found for the clones classified into these functional categories by further analyses.

35

Since the protein encoded by clones of the invention contains full-length amino acid sequence, it is possible to analyze its biological activity, and its effect on cellular conditions such as cell proliferation and differentiation by expressing the protein as a recombinant protein using an appropriate expression system, injecting the recombinant into the cell, or raising a specific antibody against the protein.

If the protein is a secretory protein, membrane protein, or protein associated with glycoprotein, signal transduction, or transcription, its biological activity can be analyzed by the methods in "Gene Transcription" (Hames B.D., and Higgins S.J. edit, (1993)), "Glycobiology" (Fukuda M., and Kobata A. edit, (1993)), "Growth Factors" (McKay I., and Leigh I. edit, (1993)), "Extracellular Matrix" (Haralson M.A., and Hassell J.R. edit, (1995), "Transcription Factors" (Latchman D.S. edit, (1993)), "Signal Transduction" (Milligans G. edit, (1992)), "Protein Phosphorylation" (Hardies G.D. edit, (1993), and "Ion Channels" (Ashley R.H. edit, (1995) featured in "The Practical Approach Series" (IRL PRESS), or "Signal Transduction Protocols"

(Kendall D.A., and Hill S.J. edit, (1995), "Glycoprotein Analysis in Biomedicine" (Hounsell E.F. edit, (1993)), featured in "Method in Molecular Biology" (Humana Press).

As to a protein associated with a disease, it is possible to perform a functional analysis as described above, but also possible to analyze correlation between the expression or the activity of the protein and a certain disease by using a specific antibody that is obtained by using expressed protein. Alternatively, it is possible to utilize the database Online Mendelian Inheritance in Man (OMIM) (http://www.ncbi.nlm.nih.gov/Omim/), which is a database of human genes and diseases, to analyze the protein.

New information is constantly being deposited in the OMIM database. Therefore, it is possible for one skilled in the art to find a new relationship between a particular disease and a gene of the present invention in the most up-to-date database.

Also, as for a secretory protein, membrane protein, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein, etc., search of the OMIM with the following keywords resulted in the finding that the proteins are associated with many diseases (the result of the OMIM search for secrete and membrane proteins is shown below).

Also, association between proteins associated to signal transduction or transcription and diseases is reported in "Transcription Factor Research-1999" (Fujii, Tamura, Kageyama, and Satake edit, (1999) Jikken-Igaku Zoukan, Vol.17, No.3), and "Gene Medicine" ((1999) Vol.3, No.2). For example, in tumors, many proteins have been shown to play a role, including secretory proteins, membrane proteins, and proteins associated with signal transduction, glycoprotein, and transcription, and also proteins associated with metabolism, cytoskeleton, and cell cycle, as described in "Tumor Biology" (Matsubara S. (1992) Syoukabou Life Science series). Thus, besides the proteins associated with diseases, many proteins described above are also potentially associated with diseases, and thus useful as a target in the medicinal industry.

The result of the OMIM search for secretory and membrane proteins is shown below, in which the keywords,

- 35 (1) secretion protein,
 - (2) membrane protein,
- 40 (3) channel, and

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(4) extracellular matrix were used.

Shown in the search result are only the accession numbers in the OMIM. Using the number, data showing the relationship between a disease and a gene or protein can be seen. The OMIM data has been renewed everyday.

1) Secretion protein

268 entries found, searching for "secretion protein"

104760, 176860, 160900, 107400, 118910, 139320, 603850, 147572,176880, 600946, 603215, 157147, 600174, 151675, 170280, 179512, 179513,138120, 179509, 246700, 179510, 600626, 179511, 600998, 109270, 601489, 154545, 179490, 185860, 603216,

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122559, 601746,147290, 602672, 146770, 603062, 179508, 131230, 601591, 602421,
       139250, 167805, 167770, 600041, 600564, 118825, 601146, 300090, 600753, 601652,
       600759, 600768, 602434, 182590, 603166, 308230, 602534, 603489, 107470, 150390,
       104610, 173120, 158106, 143890, 306900, 308700, 134797, 137350, 227500, 176300,
       107730, 600760, 138079, 120180, 120160, 120150, 124092, 138160, 101000, 227600,
       600509, 601199, 142410, 104311, 193400, 201910, 107300, 122560, 272800, 217000,
       590050, 147670, 133170, 176730, 300300, 134370, 274600, 120140,162151, 158070,
       152790, 120120, 106100, 300200, 192340, 190160, 138040, 147470, 147620, 173350,
10
       147380, 152200, 152760, 157145, 153450, 264080, 113811, 600937, 600840, 188545,
       202110, 600514, 186590, 603372, 136435, 137241, 252800, 214500, 207750, 138850,
       139191, 142640, 138130, 189907, 603692, 600633, 603355, 107270, 600377, 147892,
15
       232200, 600281, 232800, 602358, 137035, 601771, 601769, 253200, 601933, 118444,
       600270, 120700, 600945, 603732, 147660, 600761, 172400, 600823, 600877, 130080,
       171060, 107740, 307800, 602843, 130660, 152780, 124020, 601124, 601340, 601604,
       601610, 171050, 312060, 232700, 300159, 142703, 600734, 125255, 168450, 123812,
20
       188540, 147940, 188450, 600839, 182452, 188400, 182280, 176760, 263200, 600264,
       188826, 252650, 601185, 162641, 137216, 601398, 601538, 118888, 118445, 601745,
       190180, 601922, 182098, 602008, 147440, 602384, 600031, 109160, 602663, 151670,
       602682, 602730, 602779, 146880, 603061, 142704, 603140, 106150, 600732, 153620,
25
       603318, 139392, 600042, 102200, 603493, 182100, 264300, 603795, 184600
```

2) Membrane protein

1017 entries found, searching for "membrane protein"

30 130500, 305360, 153330, 173610, 170995, 109270, 170993, 309060, 120920, 602333, 133740, 133710, 602690, 133730, 159430, 600897, 133090, 601178, 602413, 602003, 109280, 603237, 602173, 107776, 602334, 125305, 602335, 182879, 154045, 309845, 600594, 603718, 603241, 603214, 603657, 603177, 600182, 601476, 602879, 136950, 35 600723, 601114, 185880, 185881, 300096, 602257, 160900, 177070, 603062, 603344, 602977, 310200, 600959, 300100, 186945, 600039, 600267, 128240, 182900, 601097, 136430, 600946, 602534, 601047, 143450, 603141, 603700, 600579, 256540, 159440, 602414, 600403, 602048, 188860, 137290, 158343, 184756, 602910, 603179, 600279, 40 108733, 107770, 173335, 602625, 154050, 219800, 603850, 601028, 600447, 104225, 186946, 601767, 603143, 121015, 603215, 227400, 603735, 600179, 602421, 180721, 176801, 176860, 600753, 603142, 176790, 600266, 601239, 115501, 143890, 121014, 121011, 125950, 603534, 304040, 601134, 600754, 601510, 601595, 190315, 300172, 45 602216, 602261, 602262, 602461, 131560, 179514, 179512, 176981, 142461, 139310, 312080, 176640, 128239, 185470, 310300, 601403, 601757, 273800, 151460, 176943, 104311, 168468, 120130, 602887, 600164, 601531, 601832,104775, 600040, 603583, .176894, 602631, 166945, 182180, 120620, 141180, 601014, 139150, 182860, 177061, 50 600174, 180069, 191275, 104760, 601693, 300017, 603518, 601009, 134651, 601107, 603868, 600168, 136425, 603531, 603291, 600917, 603216, 102720, 300118, 179590, 135630, 602285, 107450, 602296, 303630, 176878, 120090, 600322, 138160, 601212, 603293, 131230, 112205, 600763, 600718, 300187, 170715, 601966, 300051, 602474, 55 120070, 600691, 600855, 182309, 602101, 602857, 194355, 162230, 600874, 113730, 155550, 602701, 306400, 601789, 231200, 107271, 175100, 182870, 305100, 301000, 601313, 157147, 147670, 139200, 603593,157655, 600934, 155970, 602049, 155960,

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 - Channel (member of membrane protein)
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- 35 176266, 600724, 170500, 182390, 123825, 114208, 114205, 601784, 114206, 600937, 114204, 603415, 600053, 114209, 114207, 600760, 118425, 601011, 192500, 176261, 600761, 176260, 600359, 600228, 600877, 602235, 300008, 182389, 182391, 601328, 601534, 600504, 602323, 601958, 602780, 602781, 601327, 601012, 600734, 603208, 40 182392, 603220, 603219, 603888, 600054, 602232, 601745, 603537, 602604, 603796, 302910, 602866, 601013, 602905, 602906, 600163, 152427, 180901, 600702, 600308, 602754, 107776, 602024, 314555, 601949, 600235, 602023, 176263, 600681, 176265, 193245, 603305, 176258, 602983, 601219, 601141, 176267, 602343, 602726, 138253, 45 176262, 600003, 600397, 602872, 138249, 600843, 600935, 600580, 600845, 602158, 602106, 176264, 300110, 176257, 602717, 603493, 176268, 600932, 602727, 138254, 603652, 300138, 602420, 600570, 600150, 603583, 602345, 603749, 601142, 176256, 600846, 138252, 602982, 603787, 602836, 603788, 602566, 603651, 602421, 100690, 50 107777, 100725, 100710, 600509, 603061, 154275, 304040, 154276, 180902, 121014, 602368, 139311, 601383, 108745, 601313, 601042, 600131, 186360, 600109, 600229, 600170, 603319, 601485, 118503, 180903, 602076, 124030, 601059, 601212, 601218, 147450, 600855, 600919, 601154, 601157, 171060, 600968, 182139, 131230, 121015, 600421, 113730, 249210, 310500, 600637, 125950, 118800, 156490, 602974, 104610, 121011, 602522, 118504, 300041, 160900, 601382, 602103, 600465, 602014, 600442, 601109, 602481, 277900, 254210, 138247, 164920, 170280, 171050, 128100, 173910,

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4) Extracellular matrix

167 entries found, searching for "extracellular matrix"

10 603479, 602201, 601418, 601548, 154870, 115437, 602285, 602262, 602261,134797, 600754, 120361, 116935, 602263, 603320, 601807, 603321, 185250, 185261, 253700, 128239, 120324, 193300, 276901, 308700, 600514, 600261, 602109, 120140, 120150, 147557, 193400, 600536, 188826, 120180, 118661, 120320, 152200, 135821, 112260, 15 230740, 602090, 155760, 192975, 190182, 602108, 601463, 186745, 600900, 600985, 600758, 602369,179590, 601211, 600065, 602178, 600262, 182888, 182889, 151510, 182120, 150325, 190181, 150370, 186355, 193065, 165070, 154705, 147559, 146650, 146640, 153619, 175100, 187380, 231050, 188060, 135820, 156790, 130660, 301870, 20 128240, 600076, 600119, 193210, 600215, 600245, 121010, 150240, 600309, 600491, 222600, 120328, 600564, 600596, 600616, 600700, 600742, 120325, 138297, 600930. 156225, 601028, 601050, 601105, 253800, 601284, 601313, 120280, 310200, 601492, 120250, 601587, 601636, 601652, 601692, 601728, 120220, 601915, 602048, 155120, 25 310300, 120210, 120165, 120120, 118940, 116930, 602264, 116806, 602366, 120470, 602415, 602428, 602453, 602505, 602574, 603005, 603196, 603221, 603319, 107269, 216550, 103320, 603489, 603551, 603767, 603799, 603842

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There are several methods for analyzing the expression levels of genes associated with diseases. Differences in gene expression levels between diseased and normal tissues are studied by the analytical methods, for example, Northern hybridization and differential display. Other examples include a method with high-density cDNA filter, a method with DNA microarray and methods with PCR amplification (Experimental Medicine, Vol.17, No. 8, 980-1056 (1999); Cell Engineering (additional volume) DNA Microarray and Advanced PCR Methods, Muramatsu & Naba (eds.), Shujunsya). The varying levels of gene expression between diseased tussues and normal tissues can be studied by any of these analytical methods. When explicit difference in the expression level is observed for a gene, it can be conclude that the gene is closely associated with a disease or disorder. Instead of diseased tissues, cultured cells can be used for the assessment. Similarly, when gene expression is explicitly different between normal cells and cells reproducing disease-associated specific features, it can be concluded that the gene is closely associated with a disease or disorder. When the expression levels of genes are evidently varied during major cellular events (such as differentiation and apoptosis), the genes are involved in the cellular events and accordingly are candidates for disease- and/or disorderassociated genes. Further, genes exhibiting tissue-specific expression are genes playing important parts in the tissue functions and, therefore, can be candidates for genes associated with diseases and/or disorders affecting the tissues.

For example, non-enzymic protein glycation reaction is believed to be a cause for a variety of chronic diabetic complications. Accordingly, genes of which expression levels are elevated or decreased in a glycated protein-dependent manner in the endothelial cells, are associated with diabetic complications caused by glycated proteins (Diabetes 1996, 45 (Suppl. 3), S67-S72;

Diabetes, 1997, 46 (Suppl. 2), S19-S25).

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The onset of rheumatoid arthritis is thought to be involved in the proliferation of synovial cells covering inner surfaces of joint cavity and in inflammatory reaction resulted from the action of cytokines produced by leukocytes infiltrating into the joint synovial tissues (Rheumatism Information Center, http://www.rheuma-net.or.jp/). Recent studies have also revealed that tissue necrosis factor (TNF)-α participates in the onset (Current opinion in immunology 1999, 11, 657-662). When the expression of a gene exhibits responsiveness to the action of TNF on synovial cells, the gene is considered to be involved in rheumatoid arthritis. Many genes acting at the downstream of TNF-α and IL-1β among inflammation-associated cytokines have been previously identified. The respective stimulations are transduced through independent pathways of signaling cascade. There exists another signaling cascade for both stimulations, wherein NF-k B is a common transducing molecule shared by the two stimulations (J. Leukoc, Biol., 1994, 56(5): 542-547). It has also been revealed that many inflammationassociated genes, including IL-2, IL-6 and G-CSF, are varied in the expression levels thereof in response to the signal through the common pathway (Trend Genet. 1999, 15(6): 229-235). It is assumed that genes of which expression levels are varied in response to the stimulation of TNF- α or IL-1 β also participate in inflammation.

Ultraviolet radiation damage has been recognized as a risk factor for skin cancers, etc. (United States Environmental Protection Agency: Ozone Depletion Home Page, http://www.epa.gov/ozone/). Genes of which expression levels are varied in skin epidermal cells exposed to ultraviolet rays are considered to be associated with ultraviolet radiation damage of skin. In addition, genes associated with neural differentiation can be candidates for genes responsible for neurological diseases as well as candidates for genes usable for treating the diseases.

Clones exhibiting differences in the expression levels thereof can be selected by using gene expression analysis. The selection comprises, for example; analyzing cDNA clones by using high-density cDNA filter; and statistically treating the multiple signal values (signal values of radioisotope in the labeled probes or values obtained by measuring fluorescence intensities emitted from the fluorescent labels) for the respective clones by two-sample t-test, where the signal values are determined by multiple experiments of hybridization. The clones of interest are selectable based on the statistically significant differences in the signal distribution at p<0.05. However, selectable clones with significant difference in the expression levels thereof may be changed depending on the partial modification of statistical treatment. For example, the clones may be selected by conducting statistical treatment with two-sample t-test at p<0.01; or genes exhibiting more explicit differences in the expression levels thereof can be selected by performing statistical treatment with a pre-determined cut-off value for the significant signal difference. An alternative method is that the expression levels are simply compared with each other, and then, the clones of interest are selected based on the ratio of the expression levels thereof.

Clones that vary in their expression levels can also be selected by comparing the expression levels by PCR analysis, for example, by using the method of determining the band intensities representing the amounts of PCR products with ethidium bromide staining; the method of determining the values of radioisotope signals or fluorescence intensities of the PCR products when radiolabeled or fluorescent dye-labeled primers, respectively, are used in PCR amplification; or the method of determining the values of radioisotope signals or

fluorescence intensities of the probes hybridized to the PCR products when radiolabeled or fluorescent dye-labeled probes, respectively, are used in the hybridization. If the expression level ratios obtained in multiple PCR experiments are constantly at least 2-fold, such a clone can be judged to vary in its expression level. When the ratios are several-fold or not less than 10-fold, the clone can be selected as a gene exhibiting the explicit difference in its expression level.

A survey of genes of which expression levels are varied specifically to the glycated protein in the endothelial cells has revealed genes with elevated expression levels, HEMBA1003958, HEMBA1004850, MAMMA1001256, MAMMA1002132, PLACE2000411 and PLACE3000119. On the other hand, a gene of which expression level is decreased specifically to the glycated protein is MAMMA1001783. These clones are genes associated with diabetes.

A survey of genes of which expression levels are varied in response to TNF- α (Tumor Necrosis Factor-alpha) in the primary cell culture of synovial tissue has revealed the following clones with elevated expression levels in the presence of TNF- α :

HEMBA1000012, HEMBA1000005, HEMBA1000020, HEMBA1000046, HEMBA1000076, 20 HEMBA1000111, HEMBA1000168, HEMBA1000185, HEMBA1000201, HEMBA1000231. HEMBA1000243, HEMBA1000280, HEMBA1000282, HEMBA1000304, HEMBA1000307, HEMBA1000327, HEMBA1000356, HEMBA1000376, HEMBA1000387. HEMBA1000390. HEMBA1000418, HEMBA1000460, HEMBA1000491, HEMBA1000501, HEMBA1000518, 25 HEMBA1000519, HEMBA1000520, HEMBA1000531, HEMBA1000534, HEMBA1000542, HEMBA1000545, HEMBA1000591, HEMBA1000592, HEMBA1000594, HEMBA1000636, HEMBA1000655. HEMBA1000657. HEMBA1000673. HEMBA1000682. HEMBA1000686. HEMBA1000722, HEMBA1000726, HEMBA1000827, HEMBA1000870, HEMBA1000918, 30 HEMBA1000971, HEMBA1000974, HEMBA1000986, HEMBA1001019, HEMBA1001043, HEMBA1001051, HEMBA1001059, HEMBA1001060, HEMBA1001071, HEMBA1001080, HEMBA1001109, HEMBA1001140, HEMBA1001196, HEMBA1001172, HEMBA1001213, HEMBA1001226, HEMBA1001281, HEMBA1001299, HEMBA1001302, HEMBA1001303, 35 HEMBA1001323. HEMBA1001326, HEMBA1001327, HEMBA1001330, HEMBA1001351, HEMBA1001407, HEMBA1001411, HEMBA1001446, HEMBA1001454, HEMBA1001569, HEMBA1001647, HEMBA1001714, HEMBA1001804, HEMBA1001800, HEMBA1001809, HEMBA1001888, HEMBA1001912, HEMBA1001921, HEMBA1001967, HEMBA1002084, HEMBA1002161, HEMBA1002166, HEMBA1002241, HEMBA1002337, HEMBA1002363, HEMBA1002389, HEMBA1002458, HEMBA1002460, HEMBA1002469, HEMBA1002538, HEMBA1002542, HEMBA1002547, HEMBA1002609, HEMBA1002624, HEMBA1002659, HEMBA1002750, HEMBA1002770, HEMBA1002779, HEMBA1002810, HEMBA1002816, HEMBA1002818, HEMBA1002850, HEMBA1002863, HEMBA1003021, HEMBA1003033, HEMBA1003078, HEMBA1003166, HEMBA1003202, HEMBA1003204, HEMBA1003229, HEMBA1003235, HEMBA1003276, HEMBA1003286, HEMBA1003296, HEMBA1003376, HEMBA1003370, HEMBA1003403, HEMBA1003418, HEMBA1003433, HEMBA1003447, HEMBA1003560, HEMBA1003569, HEMBA1003571, HEMBA1003591, HEMBA1003597, HEMBA1003598, HEMBA1003621, HEMBA1003656. HEMBA1003662, HEMBA1003680, HEMBA1003715, HEMBA1003725, HEMBA1003729, HEMBA1003733, HEMBA1003742, HEMBA1003773, HEMBA1003783, HEMBA1003950, HEMBA1004012, HEMBA1004015, HEMBA1004048, HEMBA1004074, HEMBA1004086, HEMBA1004111, HEMBA1004131, HEMBA1004202, HEMBA1004203, HEMBA1004207, HEMBA1004248,

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	HEMBA1004274,	HEMBA1004321,	HEMBA1004330,	HEMBA1004356,	HEMBA1004366,
	HEMBA1004405,	HEMBA1004408,	HEMBA1004429,	HEMBA1004499,	HEMBA1004507,
	HEMBA1004509,	HEMBA1004542,	HEMBA1004596,	HEMBA1004604,	HEMBA1004776,
5	HEMBA1004889,	HEMBA1004934,	HEMBA1004978,	HEMBA1005019,	HEMBA1005047,
	HEMBA1005206,	HEMBA1005219,	HEMBA1005274,	HEMBA1005331,	HEMBA1005338,
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10	HEMBA1006253,	HEMBA1006268,	HEMBA1006309,	HEMBA1006377,	HEMBA1006474,
	HEMBA1006486,	HEMBA1006492,	HEMBA1006502,	HEMBA1006535,	HEMBA1006579,
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	HEMBA1007078,	HEMBA1007080,	HEMBA1007121,	HEMBA1007194,	HEMBA1007300,
15	HEMBA1007301,	HEMBA1007322,	HEMBB1000036,	HEMBB1000044,	HEMBB1000089,
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	HEMBB1000975,	HEMBB1000985,	HEMBB1001011,	HEMBB1001014,	HEMBB1001112,
20	HEMBB1001133,	HEMBB1001331,	HEMBB1001337,	HEMBB1001366,	HEMBB1001367,
	HEMBB1001384,	HEMBB1001394,	HEMBB1001429,	HEMBB1001463,	HEMBB1001619,
	HEMBB1001684,	HEMBB1001706,	HEMBB1001753,	HEMBB1001797,	HEMBB1001802,
	HEMBB1001812,	HEMBB1001874,	HEMBB1001910,	HEMBB1001915,	HEMBB1001973,
25	HEMBB1001983,	HEMBB1001990,	HEMBB1002190,	HEMBB1002193,	HEMBB1002249,
	HEMBB1002329,	HEMBB1002342,	HEMBB1002371,	HEMBB1002409,	HEMBB1002442,
	HEMBB1002489,	HEMBB1002492,	HEMBB1002520,	HEMBB1002534,	HEMBB1002596,
	HEMBB1002664,	HEMBB1002692,	HEMBB1002697,	HEMBB1002705,	MAMMA1000092,
30	MAMMA1000155,	MAMMA1000163,	MAMMA1000173,	MAMMA1000175,	MAMMA1000227,
	MAMMA1000241,	MAMMA1000257,	MAMMA1000264,	MAMMA1000266,	MAMMA1000270,
	MAMMA1000307,	MAMMA1000410,	MAMMA1000413,	MAMMA1000416,	MAMMA1000421,
	MAMMA1000472,	MAMMA1000501,	MAMMA1000605,	MAMMA1000643,	MAMMA1000670,
35	MAMMA1000684,	MAMMA1000696,	MAMMA1000732,	MAMMA1000752,	MAMMA1000802,
	MAMMA1000824,	MAMMA1000905,	MAMMA1000921,	MAMMA1000931,	MAMMA1000957,
	MAMMA1000962,				
	MAMMA1000998,	MAMMA1001008,	MAMMA1001050,	MAMMA1001074,	MAMMA1001078,
40	MAMMA1001292,	MAMMA1001397,	MAMMA1001476,	MAMMA1001743,	MAMMA1001744,
	MAMMA1001754,	MAMMA1001760,	MAMMA1001785,	MAMMA1001858,	MAMMA1001908,
	MAMMA1002236,	MAMMA1002267,	MAMMA1002292,	MAMMA1002311,	MAMMA1002322,
	MAMMA1002359,	MAMMA1002362,	MAMMA1002485,	MAMMA1002494,	MAMMA1002597,
45	MAMMA1002598,	MAMMA1002665,	MAMMA1002671,	MAMMA1002684,	MAMMA1002748,
	MAMMA1002775,	MAMMA1002830,	MAMMA1002858,	MAMMA1002868,	MAMMA1002886,
	MAMMA1002887,	MAMMA1002892,	MAMMA1002909,	MAMMA1002937,	MAMMA1002947,
	MAMMA1002964,	MAMMA1002970,	MAMMA1003013,	MAMMA1003150,	NT2RM1000039,
50	NT2RM1000062,	NT2RM1000080,	NT2RM1000086,	NT2RM1000127,	NT2RM1000132,
	NT2RM1000187,	NT2RM1000199,	NT2RM1000244,	NT2RM1000256,	NT2RM1000272,
	NT2RM1000318,	NT2RM1000354,	NT2RM1000377,	NT2RM1000430,	NT2RM1000499,
	NT2RM1000539,	NT2RM1000553,	NT2RM1000563,	NT2RM1000699,	NT2RM1000742,
55	NT2RM1000826,	NT2RM1000829,	NT2RM1000833,	NT2RM1000882,	NT2RM1000898,
	NT2RM1000905,	NT2RM1001092,	NT2RM2000013,	NT2RM2000032,	NT2RM2000042,
	NT2RM2000101,	NT2RM2000124,	NT2RM2000192,	NT2RM2000259,	NT2RM2000260,

			EI IOITOII /LE		
	NT2RM2000363,	NT2RM2000368,	NT2RM2000402,	NT2RM2000452,	NT2RM2000952,
	NT2RM2001221,	NT2RM2002014,	NT2RM2002030,	NT2RM4000156,	NT2RM4000349,
	NT2RM4000395,	NT2RM4000457,	NT2RM4000511,	NT2RM4000514,	NT2RM4000698,
5	NT2RM4000764,	NT2RM4001016, NT	2RM4001084, NT2F	RM4001594, NT2RM	4001629,
	NT2RM4001662,	NT2RM4001841,	NT2RM4002093,	NT2RM4002109,	NT2RM4002145,
	NT2RM4002189,	NT2RM4002194,	NT2RM4002226,	NT2RP1000170,	NT2RP1000439,
	NT2RP1000478,	NT2RP1000513,	NT2RP1000701,	NT2RP1000856,	NT2RP1001361,
10	NT2RP2000097,	NT2RP20002	39, NT2RP20	00288, NT2R	P2000328,
	NT2RP2000329,	NT2RP2000369,	NT2RP2000422,	NT2RP2000842,	NT2RP2000965,
	NT2RP2001245,	NT2RP2001440	NT2RP2001560,	NT2RP2001634,	NT2RP2001663,
	NT2RP2001677,	NT2RP2001762,	NT2RP2002270,	NT2RP2002312,	NT2RP2002316,
15	NT2RP2002333,	NT2RP2002706,	NT2RP2002925	NT2RP2002959,	NT2RP2002987,
	NT2RP2003125,	NT2RP2003137,	NT2RP2003237,	NT2RP2003272,	NT2RP2003596,
	NT2RP2003604,	NT2RP2003643,	NT2RP2003968,	NT2RP2003976,	NT2RP2004194,
	NT2RP2004321,	NT2RP2005037,	NT2RP2005140,	NT2RP2005204,	NT2RP2005293,
20	NT2RP2005457,	NT2RP2005555,	NT2RP2005600,	NT2RP2005701,	NT2RP2005719,
	NT2RP2005722,	NT2RP2005773,	NT2RP2005890,	NT2RP2006023,	NT2RP2006071,
	NT2RP3000186,	NT2RP3000341,	NT2RP3000599,	NT2RP3000632,	NT2RP3000644,
	NT2RP3000852,	NT2RP3000968,	NT2RP3001096,	NT2RP3001109,	NT2RP3001126,
25	NT2RP3001147,	NT2RP3001449,	NT2RP3001529	NT2RP3001753,	NT2RP3001854,
	NT2RP3001915,	NT2RP3001969,	NT2RP3002081	NT2RP3002142,	NT2RP3002399,
	NT2RP3002590,	NT2RP3002603,	NT2RP3002810,	NT2RP3002876,	NT2RP3003311.
	NT2RP3003330,	NT2RP3003672,	NT2RP3004209,	NT2RP3004378,	NT2RP4000078,
30	NT2RP4000541,	NT2RP4000588,	NT2RP4001219,	NT2RP4001228,	NT2RP4001276,
	NT2RP4001507,	NT2RP4002047,	NT2RP5003459,	NT2RP5003492,	OVARC1000085,
	OVARC1000087,	OVARC1000106	OVARC1000151,	OVARC1000198,	OVARC1000431,
	OVARC1000440	OVARC1000564,	OVARC1000605,	OVARC1000679,	OVARC1000883,
35	OVARC1000912	OVARC1000960,	OVARC1000971,	OVARC1001038,	OVARC1001055,
	OVARC1001085		OVARC1001167,	OVARC1001339,	OVARC1001425,
	OVARC1001745	·	OVARC1001766,	OVARC1001942,	OVARC1002044,
	OVARC1002138	PLACE1000004,	PLACE1000005,	PLACE1000420,	PLACE1000547,
40	PLACE1000562,	PLACE1000653,	PLACE1001168,	PLACE1001311,	PLACE1001377,
	PLACE1001920,	PLACE1001983,	PLACE1002066,	PLACE1002072,	PLACE1002140,
	PLACE1002171,	PLACE1002319,	PLACE1002474,	PLACE1002499,	PLACE1002532,
	PLACE1002665,	PLACE1003025,	PLACE1003145,	PLACE1003361,	PLACE1003605,
45	PLACE1003704,	PLACE1003783,	PLACE1003885,	PLACE1004405,	PLACE1004629,
	PLACE1004686,	PLACE1004930,	PLACE1005066,	PLACE1005077,	PLACE1005630,
	PLACE1005876,	PLACE1006143,	PLACE1006325,	PLACE1006488,	PLACE1006805,
	PLACE1006829,		PLACE1007858,	PLACE1008201,	PLACE1009045,
50	PLACE1009113,	PLACE1009621,	PLACE1010106,	PLACE1010310,	PLACE1010622,
	PLACE1010944,		PLACE1011185,	PLACE1011332,	PLACE1011635,
	PLACE1011646,	PLACE1011725,	PLACE2000014,	PLACE2000264,	PLACE2000394,
	PLACE2000419,	•	PLACE3000220,	PLACE3000254,	PLACE3000271,
55	PLACE3000339,		PLACE3000350,	PLACE3000353,	PLACE3000401,
	PLACE4000300,	•	THYRO1000855,	THYRO1001559,	Y79AA1000065,
	Y79AA1000202,	Y79AA1000214,	Y79AA1000346,	Y79AA1000784,	Y79AA1000833,
		·	•		

Y79AA1000968, Y79AA1001555, Y79AA1002220

NT2RP3000690,

THYRO1000070, THYRO1001100, Y79AA1000342

NT2RP2005358,

35

On the other hand, clones with decreased expression levels in the presence of TNF α are: NT2RM2000469, HEMBA1002150, HEMBB1000240, NT2RM2000984, NT2RM2001688, NT2RM4000290, NT2RM4000496, NT2RM4000590, NT2RM4001047, NT2RM4001582, NT2RM4001611, NT2RM4001650, NT2RM4002075, NT2RM4002128, NT2RP1000174. NT2RP1000243, NT2RP1000581, NT2RP1000688, NT2RP1000767, NT2RP1000825, NT2RP1001185; NT2RP1001286, NT2RP1001432, NT2RP1001457, NT2RP2000001. 10 NT2RP2000248, NT2RP2000841, NT2RP2001813, NT2RP2002137, NT2RP2002928, NT2RP2003517, NT2RP2003559, NT2RP2003564, NT2RP2004933, NT2RP2005038. NT2RP2006365, NT2RP3000072, NT2RP3000320, NT2RP3000484, NT2RP3000980, NT2RP3001111. NT2RP3001420, NT2RP3001495. NT2RP3002056. NT2RP3002057. 15 NT2RP3002799, NT2RP3002869, NT2RP3002953, NT2RP3002545, NT2RP3002713, NT2RP3002955. NT2RP3003282, NT2RP3003290. NT2RP3003384, NT2RP3003385, NT2RP3003870, NT2RP3004207, NT2RP3004262 NT2RP3004527, NT2RP4000500, NT2RP4000524, NT2RP4000787, NT2RP4000927, NT2RP4000955, 20 NT2RP4000989, NT2RP4001442, NT2RP4001638, NT2RP4001950, NT2RP4002888, NT2RP5003524, OVARC1001270, PLACE1000246, PLACE1002816. These are rheumatoid arthritis-associated clones. A survey of genes of which expression levels are varied in primary cultured skin fibroblast 25 cells exposed to ultraviolet light has revealed the following clones with elevated expression radiation: HEMBA1000542, by ultraviolet HEMBA1001808. HEMBA1002177. HEMBA1003314, MAMMA1001874, NT2RM2001100, NT2RP2005732, NT2RP3000592, NT2RP4000657, OVARC 1000004, OVARC1001092, OVARC1001342, PLACE1002816, 30 NT2RM4001002, NT2RM4001813, NT2RM4002266,

On the other hand, the expression levels of the following clones were decreased by ultraviolet radiation:

NT2RP3001216,

NT2RP2001174,

NT2RP3003464,

NT2RP2001196,

PLACE1006382,

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HEMBA1000005,
                    HEMBA1000150,
                                    HEMBA1000156,
                                                    HEMBA1000158,
                                                                    HEMBA1000168,
    HEMBA1000231,
                    HEMBA1000304,
                                    HEMBA1000307,
                                                    HEMBA1000333,
                                                                    HEMBA1000366,
    HEMBA1000369,
                    HEMBA1000390,
                                    HEMBA1000396,
                                                    HEMBA1000418,
                                                                    HEMBA1000434,
40
    HEMBA1000464,
                    HEMBA1000469,
                                    HEMBA1000490,
                                                    HEMBA1000504.
                                                                    HEMBA1000505.
    HEMBA1000557,
                    HEMBA1000657,
                                    HEMBA1000673,
                                                   HEMBA1000682,
                                                                    HEMBA1000686,
    HEMBA1000727,
                    HEMBA1000752,
                                    HEMBA1000851,
                                                    HEMBA1000852,
                                                                    HEMBA1000870.
    HEMBA1000872,
                    HEMBA1001085,
                                    HEMBA1001121,
                                                    HEMBA1001133,
                                                                    HEMBA1001235,
45
    HEMBA1001265,
                    HEMBA1001281,
                                    HEMBA1001289,
                                                    HEMBA1001299.
                                                                    HEMBA1001303,
    HEMBA1001310,
                    HEMBA1001323,
                                    HEMBA1001595,
                                                    HEMBA1001620,
                                                                    HEMBA1001640,
    HEMBA1001678.
                    HEMBA1001712.
                                    HEMBA1001835.
                                                    HEMBA1001950,
                                                                    HEMBA1001987,
    HEMBA1002253,
                    HEMBA1002321,
                                    HEMBA1002341,
                                                    HEMBA1002419,
                                                                    HEMBA1002679,
    HEMBA1002728,
                    HEMBA1002818,
                                    HEMBA1002935,
                                                    HEMBA1002999,
                                                                    HEMBA1003034.
                    HEMBA1003098,
    HEMBA1003071,
                                    HEMBA1003142,
                                                    HEMBA1003175,
                                                                    HEMBA1003202,
    HEMBA1003212,
                    HEMBA1003220,
                                    HEMBA1003276,
                                                    HEMBA1003373,
                                                                    HEMBA1003417,
    HEMBA1003447,
                    HEMBA1003528,
                                    HEMBA1003684,
                                                    HEMBA1003799,
                                                                    HEMBA1003885,
    HEMBA1003989.
                    HEMBA1004011.
                                    HEMBA1004055.
                                                    HEMBA1004133.
                                                                    HEMBA1004225,
    HEMBA1004272,
                                    HEMBA1004631,
                    HEMBA1004353,
                                                   HEMBA1004669,
                                                                   HEMBA1004705,
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			LI I UI 7 UII AZ		
	HEMBA1004753,	HEMBA1004776,	HEMBA1004803,	HEMBA1004816,	HEMBA1004900,
	HEMBA1005047,	HEMBA1005079,	HEMBA1005101,	HEMBA1005149,	HEMBA1005152,
	HEMBA1005202,	HEMBA1005314,	HEMBA1005372,	HEMBA1005511,	HEMBA1005513,
5	HEMBA1005518,	HEMBA1005570,	HEMBA1005577,	HEMBA1005581,	HEMBA1005588,
	HEMBA1005609,	HEMBA1005632,	HEMBA1005853,	HEMBA1006031,	HEMBA1006035,
	HEMBA1006485,	HEMBA1006486,	HEMBA1006502,	HEMBA1006696,	HEMBA1006789,
	HEMBA1006796,	HEMBA1007085,	HEMBA1007224,	HEMBA1007301,	HEMBA1007319,
10	HEMBA1007341,	HEMBA1007342,	HEMBB1000036,	HEMBB1000037,	HEMBB1000217,
	HEMBB1000266,	HEMBB1000317,	HEMBB1000336,	HEMBB1000354,	HEMBB1000369,
	HEMBB1000399,	HEMBB1000434,	HEMBB1000438,	HEMBB1000592,	HEMBB1000673,
	HEMBB1000789,	HEMBB1000810,	HEMBB1000883,	HEMBB1000887,	HEMBB1001105,
15	HEMBB1001182,	HEMBB1001242,	HEMBB1001267,	HEMBB1001424,	HEMBB1001464,
	HEMBB1001531,	HEMBB1001618,	HEMBB1001996,	HEMBB1002092,	HEMBB1002139,
	HEMBB1002142,	HEMBB1002190,	HEMBB1002453,	HEMBB1002520,	HEMBB1002550,
	HEMBB1002556,	HEMBB1002600,	HEMBB1002664,	MAMMA1000009,	MAMMA1000055,
20	MAMMA1000069,	MAMMA1000133,	MAMMA1000171,	MAMMA1000173,	MAMMA1000287,
	MAMMA1000416,	MAMMA1000585,	MAMMA1000713,	MAMMA1000760,	MAMMA1000798,
	MAMMA1000831,	MAMMA1000875,	MAMMA1000876,	MAMMA1000877,	MAMMA1000906,
	MAMMA1000931,	MAMMA1000962,	MAMMA1001133,	MAMMA1001139,	MAMMA1001243,
25	MAMMA1001271,	MAMMA1001274,	MAMMA1001298,	MAMMA1001606,	MAMMA1001630,
	MAMMA1001670,	MAMMA1001743,	MAMMA1001751,	MAMMA1002140,	MAMMA1002145,
	MAMMA1002158,	MAMMA1002170,	MAMMA1002236,	MAMMA1002311,	MAMMA1002498,
	MAMMA1002754,	MAMMA1002780,	MAMMA1002820,	MAMMA1002843,	MAMMA1002844,
30	MAMMA1002871,	MAMMA1003047,	NT2RM1000037,	NT2RM1000039,	NT2RM1000080,
	NT2RM1000086,	NT2RM1000341,	NT2RM1000499,	NT2RM1000669,	NT2RM1000746,
	NT2RM1000781,	NT2RM1000885,	NT2RM1000905,	NT2RM1000962,	NT2RM2000239,
	NT2RM2000260,	NT2RM2000371,	NT2RM2000639,	NT2RM2000649,	NT2RM2000735,
35	NT2RM2000821,	NT2RM2000984,	NT2RM2001035,	NT2RM2001065,	NT2RM2001105,
	NT2RM2001177,	NT2RM2001194,	NT2RM2001196,	NT2RM2001243,	NT2RM2001256,
	NT2RM2001424,	NT2RM2001588,	NT2RM2001635,	NT2RM2001648,	NT2RM2001652,
	NT2RM2001668,	NT2RM2001706,	NT2RM2001727,	NT2RM2001730,	NT2RM2001743,
40	NT2RM2001753,	NT2RM2001760,	NT2RM2001771,	NT2RM2001785,	NT2RM2001800,
	NT2RM2001855,	NT2RM2001896,	NT2RM2001997,	NT2RM2002030,	NT2RM2002049,
	NT2RM2002091,	NT2RM2002142,	NT2RM2002145,	NT2RM2002178,	NT2RM2002580,
45	NT2RM4000215,	NT2RM4000344,	NT2RM4000368,	NT2RM4000421,	NT2RM4000425,
45	NT2RM4000457,	NT2RM4000496,	NT2RM4000515,	NT2RM4000712,	NT2RM4000787,
	NT2RM4000813,	NT2RM4000820,	NT2RM4000852,	NT2RM4000950,	NT2RM4000996,
	NT2RM4001016,	NT2RM4001047,	NT2RM4001054,	NT2RM4001140,	NT2RM4001151,
50	NT2RM4001187,	NT2RM4001204,	NT2RM4001258,	NT2RM4001437,	NT2RM4001454,
50	NT2RM4001489,	NT2RM4001605,	NT2RM4001611,	NT2RM4001666,	NT2RM4001710,
	NT2RM4001714,	NT2RM4001715,	NT2RM4001731,	NT2RM4001741,	NT2RM4001746,
	NT2RM4001856,	NT2RM4001938,	NT2RM4001940,	NT2RM4001984,	NT2RM4001987,
55	NT2RM4002013,	NT2RM4002054,	NT2RM4002055,	NT2RM4002073,	NT2RM4002145,
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	NT2RM4002213,	NT2RM4002266,	NT2RM4002278,	NT2RM4002558,	NT2RM4002565,
	NT2RM4002593,	NT2RM4002594,	NT2RP1000063,	NT2RP1000111,	NT2RP1000124,

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	NT2RP1000174,	NT2RP1000348,	NT2RP1000443,	NT2RP1000522,	NT2RP1000677,
	NT2RP1000695,	NT2RP1000730,	NT2RP1000767,	NT2RP1000834,	NT2RP1000856,
	NT2RP1000944,	NT2RP1001014,	NT2RP1001073,	NT2RP1001079,	NT2RP1001177,
5	NT2RP1001185,	NT2RP1001199,	NT2RP1001253,	NT2RP1001310,	NT2RP1001313,
	NT2RP1001385,	NT2RP1001410,	NT2RP1001449,	NT2RP1001546,	NT2RP2000120,
	NT2RP2000126,	NT2RP2000205,	NT2RP2000224,	NT2RP2000274,	NT2RP2000298,
	NT2RP2000310,	NT2RP2000327,	NT2RP2000328,	NT2RP2000329,	NT2RP2000337,
10	NT2RP2000369,	NT2RP2000414,	NT2RP2000498,	NT2RP2000510,	NT2RP2000715,
	NT2RP2000819,	NT2RP2000931,	NT2RP2000938,	NT2RP2001044,	NT2RP2001065,
	NT2RP2001081,	NT2RP2001137,	NT2RP2001312,	NT2RP2001327,	NT2RP2001366,
	NT2RP2001381,	NT2RP2001420,	NT2RP2001427,	NT2RP2001449,	NT2RP2001511,
15	NT2RP2001526,	NT2RP2001560,	NT2RP2001576,	NT2RP2001581,	NT2RP2001601,
	NT2RP2001613,	NT2RP2001628,	NT2RP2001663,	NT2RP2001675,	NT2RP2001677,
	NT2RP2001678,	NT2RP2001699,	NT2RP2001721,	NT2RP2001740,	NT2RP2001813,
	NT2RP2001861,	NT2RP2001876,	NT2RP2001907,	NT2RP2001926,	NT2RP2001946,
20	NT2RP2001985,	NT2RP2002032,	NT2RP2002046,	NT2RP2002076,	NT2RP2002099,
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25	NT2RP2002710,	NT2RP2002750,	NT2RP2002800,	NT2RP2002862,	NT2RP2002880,
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	NT2RP2004081,	NT2RP2004142,	NT2RP2004165,	NT2RP2004170,	NT2RP2004194,
35	NT2RP2004196,	NT2RP2004239,	NT2RP2004240	,NT2RP2004321,	NT2RP2004364,
	NT2RP2004365,	NT2RP2004373,	NT2RP2004389,	NT2RP2004392,	NT2RP2004399,
	NT2RP2004538,	NT2RP2004587,	NT2RP2004600,	NT2RP2004602,	NT2RP2004614,
40	NT2RP2004664,	NT2RP2004675,	NT2RP2004681,	NT2RP2004710,	NT2RP2004743,
40	NT2RP2004767,	NT2RP2004799,	NT2RP2004816,	NT2RP2004959,	NT2RP2004962,
	NT2RP2005000,	NT2RP2005003,	NT2RP2005012,	NT2RP2005031,	NT2RP2005037,
	NT2RP2005108,	NT2RP2005126,	NT2RP2005144,	NT2RP2005147,	NT2RP2005289,
45	NT2RP2005293,	NT2RP2005325,	NT2RP2005354,	NT2RP2005441,	NT2RP2005464,
40	NT2RP2005491,	NT2RP2005495,	NT2RP2005498,	NT2RP2005501,	NT2RP2005555,
	NT2RP2005557,	NT2RP2005605,	NT2RP2005622,	NT2RP2005645,	NT2RP2005651,
	NT2RP2005654,	NT2RP2005669,	NT2RP2005694,	NT2RP2005701,	NT2RP2005752,
50	NT2RP2005773,	NT2RP2005804,	NT2RP2005835,	NT2RP2005841,	NT2RP2005857,
••	NT2RP2005890,	NT2RP2005901,	NT2RP2005942,	NT2RP2006023,	NT2RP2006052,
	NT2RP2006069,	NT2RP2006166,	NT2RP2006184,	NT2RP2006186,	NT2RP2006196,
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40				8, MAMMA1001080,	
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15	THYRO1000197,	THYRO1000221,	THYRO1000253,	THYRO1000270,	THYRO1000279,
	THYRO1000327,	THYRO1000394,	THYRO1000438,	THYRO1000558,	THYRO1000569,
	THYRO1000585,	THYRO1000596,	THYRO1000625,	THYRO1000637,	THYRO1000676,
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	THYRO1001772,	THYRO1001854,	VESEN1000122,	Y79AA1000037,	Y79AA1000065,
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25	Y79AA1000538,	Y79AA1000782,	Y79AA1001023,	Y79AA1001145,	Y79AA1001391,
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	Y79AA1001963, Y	79AA1002125, Y79A	A1002229, Y79AA10	02407, Y79AA10024	187

These clones are associated with ultraviolet radiation damage.

A survey of genes of which expression levels are varied in response to the stimulation for inducing cell differentiation (stimulation using retinoic acid (RA) or using RA/inhibitor (inhibitor for cell division)) in culture cells of neural strain, NT2, revealed the following clones with elevated expression levels in the presence of RA:

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                                                  HEMBA1000376,
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                                                                        Y79AA1002407,
    Y79AA1002472, Y79AA1002482
      On the other hand, clones of which expression levels decreased by RA are as follows:
    HEMBA1000946,
                     HEMBA1003569,
                                      HEMBA1005570,
                                                      HEMBB1000915,
                                                                       NT2RM1000666,
    NT2RM2000092,
                     NT2RM2000594.
                                      NT2RM2001256,
                                                      NT2RM4001754,
                                                                       NT2RM4001905,
    NT2RP2001675,
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                                      NT2RP2005491,
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                                                                       NT2RP3002081,
    NT2RP3004594,
                     NT2RP4001950,
                                     NT2RP4002408,
                                                      OVARC1000431,
                                                                       OVARC1001942,
    OVARC1001943,
                      PLACE1003190,
                                      PLACE1004868,
                                                      PLACE1005923,
                                                                       PLACE1007257,
    PLACE1010624, Y79AA1000346
      Clones of which expression levels increase by RAInhibitor are as follows:
    HEMBA1000046,
                     HEMBA1000307,
                                      HEMBA1000434,
                                                      HEMBA1000504,
                                                                       HEMBA1000588,
45
    HEMBA1000682,
                     HEMBA1000726.
                                      HEMBA1000943.
                                                      HEMBA1001071.
                                                                       HEMBA1001094.
    HEMBA1001122,
                     HEMBA1001323,
                                      HEMBA1001361,
                                                      HEMBA1001455,
                                                                       HEMBA1001709,
    HEMBA1001746,
                     HEMBA1001869,
                                      HEMBA1002084,
                                                      HEMBA1002583,
                                                                       HEMBA1002628,
    HEMBA1002801,
                     HEMBA1002937,
                                      HEMBA1003096,
                                                      HEMBA1003142,
                                                                       HEMBA1003229,
    HEMBA1003276,
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                                                      HEMBA1003597,
                                                                       HEMBA1003617,
    HEMBA1003725,
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                                      HEMBA1003879,
                                                      HEMBA1003989,
                                                                       HEMBA1004000,
    HEMBA1004015,
                     HEMBA1004024
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5	HEMBA1006364,	HEMBA1006485,	HEMBA1006559,	HEMBA1006579,	HEMBA1006754,
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	HEMBB1000852,	HEMBB1000913,	HEMBB1000985,	HEMBB1001117,	HEMBB1001210,
	HEMBB1001317,	HEMBB1001394,	HEMBB1001443,	HEMBB1001668,	HEMBB1001695,
10	HEMBB1002049,	HEMBB1002254,	HEMBB1002266,	HEMBB1002371,	HEMBB1002502,
	HEMBB1002614,	HEMBB1002617,	HEMBB1002692,	HEMBB1002697,	MAMMA1000241,
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15	MAMMA1001820, I	MAMMA1002267, MA	MMA1002769, MAM	MA1002871, MAMM/	A1002941,
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	NT2RP2006441,	NT2RP2006467,	NT2RP2006534,	NT2RP2006565,	NT2RP3000348,
50	NT2RP3000359,	NT2RP3000366,	NT2RP3000403,	NT2RP3000418,	NT2RP3000441,
	NT2RP3000561,	NT2RP3000759,	NT2RP3000826,	NT2RP3001007,	NT2RP3001096,
	NT2RP3001126,	NT2RP3001355,	NT2RP3001396,	NT2RP3001449,	NT2RP3001490,
	NT2RP3001679,	NT2RP3001727,	NT2RP3001752,	NT2RP3001777,	NT2RP3001782,
55	NT2RP3001799,	NT2RP3001854,	NT2RP3001989,	NT2RP3002142,	NT2RP3002248,
	NT2RP3002343,	NT2RP3002484,	NT2RP3002529,	NT2RP3002549,	NT2RP3002628,
	NT2RP3002687,	NT2RP3002688,	NT2RP3002810,	NT2RP3003032,	NT2RP3003139,

			LI I UI W U II AZ		
	NT2RP3003193,	NT2RP3003203,	NT2RP3003204,	NT2RP3003210,	NT2RP3003212,
	NT2RP3003264,	NT2RP3003282,	NT2RP3003500,	NT2RP3004041,	NT2RP3004215,
	NT2RP4000147,	NT2RP4000259,	NT2RP4000360,	NT2RP4000448,	NT2RP4000524,
5	NT2RP4000588,	NT2RP4000879,	NT2RP4000907,	NT2RP4000989,	NT2RP4001079,
	NT2RP4001150,	NT2RP4001219,	NT2RP4001260,	NT2RP4001274,	NT2RP4001353,
	NT2RP4001547,	NT2RP4001677,	NT2RP4002052,	OVARC1000006,	OVARC1000092,
	OVARC1000321,	OVARC1000384,	OVARC1000408,	OVARC1000414,	OVARC1000520,
10	OVARC1000526,	OVARC1000588,	OVARC1000679,	OVARC1000682,	OVARC1000769,
	OVARC1000850,	OVARC1000862,	OVARC1000886,	OVARC1000984,	OVARC1001000,
	OVARC1001004,	OVARC1001154,	OVARC1001170,	OVARC1001173,	OVARC1001200,
	OVARC1001268,	OVARC1001376,	OVARC1001419,	OVARC1001425,	OVARC1001476,
15	OVARC1001480,	OVARC1001542,	OVARC1001873,	OVARC1001928,	OVARC1001987,
	OVARC1002066,	OVARC1002082, OV	'ARC1002112, OVAI	RC1002127,	
	PLACE1000014,	PLACE1000048,	PLACE1000184,	PLACE1000185,	PLACE1000246,
	PLACE1000292,	PLACE1000332,	PLACE1000347,	PLACE1000564,	PLACE1000656,
20	PLACE1000712,	PLACE1001000,	PLACE1001168,	PLACE1001185,	PLACE1001241,
	PLACE1001294,	PLACE1001311,	PLACE1001395,	PLACE1001570,	PLACE1001608,
	PLACE1001610,	PLACE1001716,	PLACE1001746,	PLACE1001817,	PLACE1001821,
	PLACE1001844,	PLACE1001897,	PLACE1002066,	PLACE1002119,	PLACE1002157,
25	PLACE1002205,	PLACE1002256,	PLACE1002259,	PLACE1002399,	PLACE1002438,
	PLACE1002474,	PLACE100247	77, PLACE10	02500, PLAC	E1002514,
	PLACE1002578,	PLACE1002815,	PLACE1002851,	PLACE1002968,	PLACE1003108,
	PLACE1003174,	PLACE1003200,	PLACE1003238,	PLACE1003256,	PLACE1003334,
30	PLACE1003342,	PLACE1003516,	PLACE1003521,	PLACE1003537,	PLACE1003592,
	PLACE1003596,	PLACE1003723,	PLACE1003760,	PLACE1003771,	PLACE1003783,
	PLACE1003795,	PLACE1003892,	PLACE1003968,	PLACE1004103,	PLACE1004256,
	PLACE1004405,	PLACE1004460,	PLACE1004506,	PLACE1004629,	PLACE1004674,
35	PLACE1004813,	PLACE1004979,	PLACE1005066,	PLACE1005101,	PLACE1005102,
	PLACE1005128,	PLACE1005181,	PLACE1005287,	PLACE1005305,	PLACE1005327,
	PLACE1005477,	PLACE1005595,	PLACE1005603,	PLACE1005666,	PLACE1005804,
40	PLACE1005884,	PLACE1005934,	PLACE1006076,	PLACE1006119,	PLACE1006159,
40	PLACE1006164,	PLACE1006170,	PLACE1006382,	PLACE1006492,	PLACE1006629,
	PLACE1006704,	PLACE1006731,	PLACE1006760,	PLACE1006779,	PLACE1006795,
	PLACE1006805,	PLACE1006962,	PLACE1007045,	PLACE1007111,	PLACE1007282,
45	PLACE1007386,	PLACE1007416,	PLACE1007484,	PLACE1007544,	PLACE1007645,
45	PLACE1007743,	PLACE1007746,	PLACE1007807,	PLACE1007858,	PLACE1008002,
	PLACE1008181,	PLACE1008273,	PLACE1008368,	PLACE1008405,	PLACE1008532,
	PLACE1008568,	PLACE1008625,	PLACE1008696,	PLACE1008867,	PLACE1009027,
50	PLACE1009039,	PLACE1009045,	PLACE1009110,	PLACE1009298,	PLACE1009328,
50	PLACE1009581,	PLACE1009621,	PLACE1009622,	PLACE1009637,	PLACE1009925,
	PLACE1009935,	PLACE1010089,	PLACE1010106,	PLACE1010152,	PLACE1010274,
	PLACE1010491,	PLACE1010629,	PLACE1010630,	PLACE1010714,	PLACE1010739,
55	PLACE1010891,	PLACE1010896,	PLACE1010925,	PLACE1010965,	PLACE1011026,
-	PLACE1011046,	PLACE1011214,	PLACE1011399,	PLACE1011433,	PLACE1011492,
	PLACE1011641,	PLACE1011649,	PLACE1011719,	PLACE1011762,	PLACE1011858,
	PLACE1011923,	PLACE2000014,	PLACE2000039,	PLACE2000216,	PLACE2000302,

			LI I VI V VII AL		
	PLACE2000317,	PLACE2000342,	PLACE2000347,	PLACE2000379,	PLACE3000121,
	PLACE3000124,	PLACE3000160,	PLACE3000242,	PLACE3000271,	PLACE3000353,
	PLACE3000362,	PLACE3000365,	PLACE3000400,	PLACE3000401,	PLACE4000034,
5	PLACE4000089, F	PLACE4000522, PLA	ACE4000558,		
	SKNMC1000050,	THYRO1000040,	THYRO1000197,	THYRO1000241,	THYRO1000327,
	THYRO1000394,	THYRO1000488,	THYRO1000501,	THYRO1000585,	THYRO1000596,
	THYRO1000625,	THYRO1000805,	THYRO1000934,	THYRO1001133,	THYRO1001134,
10	THYRO1001173,	THYRO1001213,	THYRO1001262,	THYRO1001290,	THYRO1001721,
	Y79AA1000037,	Y79AA1000800,	Y79AA1000976,	Y79AA1001078,	Y79AA1001228,
	Y79AA1001299,	Y79AA1001402,	Y79AA1001585,	Y79AA1001696,	Y79AA1001711,
	Y79AA1001827,	Y79AA1001875,	Y79AA1002027,	Y79AA1002211,	Y79AA1002234,
15	Y79AA1002258				
	On the other ha	and, clones of which	expression levels de	crease by RA/inhibite	or are as follows:
	HEMBA1000012,	HEMBA1000501,	HEMBA1000946,	HEMBA1003220,	HEMBA1003403,
	HEMBA1003569,	HEMBA1003591,	HEMBA1003926,	HEMBA1004168,	HEMBA1004507,
20	HEMBA1005009,	HEMBA1005296,	HEMBA1005528,	HEMBA1005570,	HEMBA1006467,
	HEMBA1006486,	HEMBA1006492,	HEMBA1007322,	HEMBB1000055,	HEMBB1000244,
	HEMBB1001665,	MAMMA1000684,	MAMMA1001139,	MAMMA1001743,	NT2RM1000257
	NT2RM1000318,	NT2RM1000539,	NT2RM1000666,	NT2RM2000092,	NT2RM2000192,
25	NT2RM2000371,	NT2RM2000594,	NT2RM4000511,	NT2RM4001140,	NT2RM4001754,
	NT2RM4001905,	NT2RM4001940,	NT2RM4002593,	NT2RP1000086,	NT2RP1000439,
	NT2RP1001073,	NT2RP2000098,	NT2RP2000965,	NT2RP2001397,	NT2RP2002047,
	NT2RP2004226,	NT2RP2004396,	NT2RP2004655,	NT2RP2005126,	NT2RP2005464,
30	NT2RP2005712,	NT2RP2005859,	NT2RP2005890,	NT2RP3000980,	NT2RP3001383,
	NT2RP3001621,	NT2RP3002081,	NT2RP3002181,	NT2RP3002244,	NT2RP3002590,
	NT2RP3003059,	NT2RP300425	58, NT2RP300	04378, NT2RI	P3004527,
	NT2RP3004594,	NT2RP4001760,	NT2RP4001950,	NT2RP4002047,	NT2RP4002408,
35	NT2RP5003459,	OVARC1000004,	OVARC1000035,	OVARC1000431,	OVARC1001051,
	OVARC1001129,	OVARC1001176,	OVARC1001261,	OVARC1001342,	OVARC1001942,
	OVARC1001943,	PLACE1002171,	PLACE1002465,	PLACE1003190,	PLACE1003375,
	PLACE1004128,	PLACE1005026,	PLACE1005876,	PLACE1005923,	PLACE1007257,
40	PLACE1007375,	PLACE1007507,	PLACE1008941,	PLACE1010624,	PLACE1011090,
	PLACE1011219, T	HYRO1000270, Y79	AA1000346, Y79AA	1001541	

These clones are also associated with neural differentiation and, therefore, are candidates for genes associated with neurological diseases.

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For example, if the protein encoded by the cDNA of the present invention is a regulatory factor of cellular conditions such as growth and differentiation, it can be used for developing medicines as follows. The protein or antibody provided by the invention is injected into a certain kind of cells by microinjection. Then, using the cells, it is possible to screen low molecular weight compounds by measuring the change in the cellular conditions, or the activation or inhibition of a particular gene. The screening can be performed as follows. First, the protein is expressed and purified as recombinant. The purified protein is microinjected into cells such as various cell lines, or primary culture cells, and the cellular change such as growth and differentiation can be examined. Alternatively, the induction of genes whose expression is known to be associated with a particular change of cellular conditions may be

detected by the amount of mRNA or protein. Or, the amount of intracellular molecules (low molecular weight compounds, etc.) that is changed by the function of the gene product (protein) which is known to be associated with a particular change of cellular conditions may be detected. The compounds to be screened (both low and high molecular compounds are acceptable) can be added to the culture media and assessed for their activity by measuring the change of the cellular conditions. Instead of microinjection, cell lines introduced with the gene obtained in the invention can be used for the screening. If the gene product is turn out to be associated with a particular change in the cellular conditions, the change of the product can be used as a measurement for screening. Once a compound is screened out which can activate or inhibit the function of the protein of the invention, it can be applied for developing medicines.

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If the protein encoded by the cDNA of the present invention is a secretory protein, membrane protein, or protein associated with signal transduction, glycoprotein, transcription, or diseases, it can be used in functional assays for developing medicines.

In case of a membrane protein, it is most likely to be a protein that functions as a receptor or ligand on the cell surface. Therefore, it is possible to reveal a new relationship between a ligand and receptor by screening the membrane protein of the invention based on the binding activity with the known ligand or receptor. Screening can be performed according to the known methods.

For example, a ligand against the protein of the invention can be screened in the following manner. Namely, a ligand that binds to a specific protein can be screened by a method comprising the steps of: (a) contacting a test sample with the protein of the invention or a partial peptide thereof, or cells expressing these, and (b) selecting a test sample that binds to said protein, said partial peptide, or said cells.

On the other hand, for example, screening using cells expressing the protein of the present invention that is a receptor protein can also be performed as follows. It is possible to screen receptors that is capable of binding to a specific protein by using procedures (a) attaching the sample cells to the protein of the invention or its partial peptide, and (b) selecting cells that can bind to the said protein or its partial peptide.

In a following screening as an example, first the protein of the invention is expressed, and the recombinant protein is purified. Next, the purified protein is labeled, binding assay is performed using a various cell lines or primary cultured cells, and cells that are expressing a receptor are selected (Growth and differentiation factors and their receptors, Shin-Seikagaku Jikken Kouza Vol.7 (1991) Honjyo, Arai, Taniguchi, and Muramatsu edit, p203-236, Tokyo-Kagaku-Doujin). A protein of the invention can be labeled with RI such as ¹²⁵al, and enzyme (alkaline phosphatase etc.). Alternatively, a protein of the invention may be used without labeling and then detected by using a labeled antibody against the protein. The cells that are selected by the above screening methods, which express a receptor of the protein of the invention, can be used for the further screening of an agonists or antagonists of the said receptor.

Once the ligand binding to the protein of the invention, the receptor of the protein of the invention or the cells expressing the receptor are obtained by screening, it is possible to screen a compound that binds to the ligand and receptor. Also it is possible to screen a compound that can inhibit both bindings (agonists or antagonists of the receptor, for example) by utilizing the binding activities.

When the protein of the invention is a receptor, the screening method comprises the steps of

(a) contacting the protein of the invention or cells expressing the protein of the invention with the ligand, in the presence of a test sample, (b) detecting the binding activity between said protein or cells expressing said protein and the ligand, and (c) selecting a compound that reduces said binding activity when compared to the activity in the absence of the test sample. Furthermore, when the protein of the invention is a ligand, the screening method comprises the steps of (a) contacting the protein of the invention with its receptor or cells expressing the receptor in the presence of samples, (b) detecting the binding activity between the protein and its receptor or the cells expressing the receptor, and (c) selecting a compound that can potentially reduce the binding activity compared to the activity in the absence of the sample.

Samples to screen include cell extracts, expressed products from a gene library, synthesized low molecular compound, synthesized peptide, and natural compounds, for example, but are not construed to be listed here. A compound that is isolated by the above screening using a binding activity of the protein of the invention can also be used as a sample.

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A compound isolated by the screening may be a candidate to be an agonist or an antagonist of the receptor of the protein. By utilizing an assay that monitors a change in the intracellular signaling such as phosphorylation which results from reduction of the binding between the protein and its receptor, it is possible to identify whether the obtained compound is an agonist or antagonist of the receptor. Also, the compound may be a candidate of a molecule that can inhibit the interaction between the protein and its associated proteins (including a receptor) in vivo. Such compounds can be used for developing drugs for precaution or cures of a disease with which the protein is associated.

Secretory proteins may regulate cellular conditions such as growth and differentiation. It is possible to find out a novel factor that regulates cellular conditions by adding the secretory protein of the invention to a certain kind of cell, and performing a screening by utilizing the cellular changes in growth or differentiation, or activation of a particular gene.

The screening can be performed, for example, as follows. First, the protein of the invention is expressed and purified in a recombinant form. Then, the purified protein is added to a various kind of cell lines or primary cultured cells, and the change in the cell growth and differentiation is monitored. The induction of a particular gene that is known to be involved in a certain cellular change is detected by the amounts of mRNA and protein. Alternatively, the amount of an intracellular molecule (low-molecular-weight compounds, etc.) that is changed by the function of a gene product (protein) that is known to function in a certain cellular change is used for the detection.

Once the screening reveals that the protein of the invention can regulate cellular conditions or the functions, it is possible to apply the protein as a pharmaceutical and diagnostic medicine for associated diseases by itself or by altering a part of it into an appropriate composition.

As is above described for membrane proteins, the secretory protein provided by the invention may be used to explore a novel ligand-receptor interaction using a screening based on the binding activity to a known ligand or receptor. A similar method can be used to identify an agonist or antagonist. The resulting compounds obtained by the methods can be a candidate of a compound that can inhibit the interaction between the protein of the invention and an interacting molecule (including a receptor). The compounds may be able to use as a preventive, therapeutic, and diagnostic medicine for the diseases, in which the protein may play a certain role.

Proteins associated with signal transduction or transcription may be a factor that affects a certain protein or gene in response to intracellular/extracellular stimuli. It is possible to find out a novel factor that can affect a protein or gene by expressing the protein provided by the invention in a certain types of cells, and performing a screening utilizing the activation of a certain intracellular protein or gene.

The screening may be performed as follows. First, a transformed cell line expressing the protein is obtained. Then, the transformed cell line and the untransformed original cell line are compared for the changes in the expression of a certain gene by detecting the amount of its mRNA or protein. Alternatively, the amount of an intracellular molecule (low molecular weight compounds) that is changed by the function of a certain gene product (protein) may be used for the detection. Furthermore, the change of the expression of a certain gene can be detected by introducing a fusion gene that comprises a regulatory region of the gene and a marker gene (luciferase, beta-galactosidase, etc.) into a cell, expressing the protein provided by the invention into the cell, and estimating the activity of a marker gene product (protein).

If the protein or gene of the invention is associated with diseases, it is possible to screen a gene or compound that can regulate its expression and/or activity either directly or indirectly by utilizing the protein of the present invention.

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For example, the protein of the invention is expressed and purified as a recombinant protein. Then, the protein or gene that interacts with the protein of the invention is purified, and screened based on the binding. Alternatively, the screening can be performed by adding with a compound of a candidate of the inhibitor added in advance and monitoring the change of binding activity. In another method, a transcription regulatory region locating in the 5'-upstream of the gene encoding the protein of the invention that is capable of regulating the expression of other genes is obtained, and fused with a marker gene. The fusion is introduced into a cell, and the cell is added with compounds to explore a regulatory factor of the expression of the said gene.

The compound obtained by the screening can be used for developing pharmaceutical and diagnostic medicines for the diseases with which the protein of the present invention is associated. Similarly, if the regulatory factor obtained in the screening is turn out to be a protein, compounds that can newly affect the expression or activity of the protein may be used as a medicine for the diseases with which the protein of the invention is associated.

If the protein of the invention has an enzymatic activity, regardless as to whether it is a secretory protein, membrane protein, or proteins associated with signal transduction, glycoprotein, transcription, or diseases, a screening may be performed by adding a compound to the protein of the invention and monitoring the change of the compound. The enzymatic activity may also be utilized to screen a compound that can inhibit the activity of the protein.

In a screening given as an example, the protein of the invention is expressed and the recombinant protein is purified. Then, compounds are contacted with the purified protein, and the amount of the compound and the reaction products is examined. Alternatively, compounds that are candidates of an inhibitor are pretreated, then a compound (substrate) that can react with the purified protein is added, and the amount of the substrate and the reaction products is examined.

The compounds obtained in the screening may be used as a medicine for diseases with which the protein of the invention is associated. Also they can be applied for tests that examine whether the protein of the invention functions normally *in vivo*.

Whether the secretory protein, membrane protein, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein of the present invention is a novel protein associated with diseases or not is determined in another method than described above, by obtaining a specific antibody against the protein of the invention, and examining the relationship between the expression or activity of the protein and a certain disease. In an alternative way, it may be analyzed referred to the methods in "Molecular Diagnosis of Genetic Diseases" (Elles R. edit, (1996) in the series of "Method in Molecular Biology" (Humana Press).

Proteins associated with diseases are targets of screening as mentioned, and thus are very useful in developing drugs which regulate their expression and activity. Also, the proteins are useful in the medicinal industry as a diagnostic marker of the associated disease or a target of gene therapy.

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Compounds isolated as mentioned above can be administered patients as it is, or after formulated into a pharmaceutical composition according to the known methods. For example, a pharmaceutically acceptable carrier or vehicle, specifically sterilized water, saline, plant oil, emulsifier, or suspending agent can be mixed with the compounds appropriately. The pharmaceutical compositions can be administered to patients by a method known to those skilled in the art, such as intraarterial, intravenous, or subcutaneous injections. The dosage may vary depending on the weight or age of a patient, or the method of administration, but those skilled in the art can choose an appropriate dosage properly. If the compound is encoded by DNA, the DNA can be cloned into a vector for gene therapy, and used for gene therapy. The dosage of the DNA and the method of its administration may vary depending on the weight or age of a patient, or the symptoms, but those skilled in the art can choose properly.

The present invention further relates to databases comprising at least a sequence of polynucleotide and/or protein, or a medium recorded in such databases, selected from the sequence data of the nucleotide and/or the amino acids indicated in Table 350 and Table 351. The term "database" means a set of accumulated information as machine-searchable and readable information of nucleotide sequence. The databases of the present invention comprise at least one of the novel nucleotide sequences of polynucleotides provided by the present invention. The databases of the present invention can consist of only the sequence data of the novel polynucleotides provided by the present invention or can comprise other information on nucleotide sequences of known full-length cDNAs or ESTs. The databases of the present invention can be comprised of not only the information on the nucleotide sequences but also the information on the gene functions revealed by the present invention. Additional information such as names of DNA clones carrying the full-length cDNAs can be recorded or linked together with the sequence data in the databases.

The database of the present invention is useful for gaining complete gene sequence information from partial sequence information of a gene of interest. The database of the present invention comprises nucleotide sequence information of full-length cDNAs. Consequently, by comparing the information in this database with the nucleotide sequence of a partial gene fragment yielded by differential display method or subtraction method, the information on the full-length nucleotide sequence of interest can be gained from the sequence of the partial fragment as a starting clue.

The sequence information of the full-length cDNAs constituting the database of the present invention contains not only the information on the complete sequences but also extra information on expression frequency of the genes as well as homology of the genes to known genes and

known proteins. Thus the extra information facilitates rapid functional analyses of partial gene fragments. Further, the information on human genes is accumulated in the database of the present invention, and therefore, the database is useful for isolating a human homologue of a gene originating from other species. The human homologue can be isolated based on the nucleotide sequence of the gene from the original species.

At present, information on a wide variety of gene fragments can be obtained by differential display method and subtraction method. In general, these gene fragments are utilized as tools for isolating the full-length sequences thereof. When the gene fragment corresponds to an already-known gene, the full-length sequence is easily obtained by comparing the partial sequence with the information in known databases. However, when there exists no information corresponding to the partial sequence of interest in the known databases, cDNA cloning should be carried out for the full-length cDNA. It is often difficult to obtain the full-length nucleotide sequence using the partial sequence information as an initial clue. If the full-length of the gene is not available, the amino acid sequence of the protein encoded by the gene remains unidentified. Thus the database of the present invention can contribute to the identification of full-length cDNAs corresponding to gene fragments, which cannot be revealed by using databases of known genes.

The present invention has provided 5602 novel full-length cDNA clones, and primers for synthesizing the cDNA. As has not yet proceeded the isolation of full-length cDNA within the human, the invention has great significance. The full-length cDNA clones contain the translation initiation site, and thus provide a useful information for analysis of protein functions.

The cDNA clones are assumed to encode proteins such as secretory proteins, membrane proteins, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein, etc., which have important functions in vivo, and also predicted to be associated with many diseases. The genes and proteins associated with diseases are useful for developing a diagnostic marker or medicines for regulation of their expression and activity, or as a target of gene therapy.

The invention is illustrated more specifically with reference to the following examples, but is not to be construed as being limited thereto.

EXAMPLE 1

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Construction of a cDNA library by the oligo-capping method.

The NT-2 neuron progenitor cells (Stratagene), a teratocarcinoma cell line from human embryo testis, which can differentiate into neurons by the treatment with retinoic acid were used. The NT-2 cells were cultured according to the manufacturer's instructions as follows.

- (1) NT-2 cells were cultured without induction by retinoic acid treatment (NT2RM1, NT2RM2, NT2RM4).
 - (2) After cultured, NT-2 cells were induced by adding retinoic acid, and then were cultured for 48 hours (NT2RP1).
 - (3) After cultured, NT-2 cells were induced by adding retinoic acid, and then were cultured

for 2 weeks (NT2RP2, NT2RP3, NT2RP4, NT2RP5).

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Also, the human neuroblastoma cell line SK-N-MC (ATCC HTB-10) (SKNMC1), and human retinoblastoma cell line Y79 (ATCC HTB-18) (Y79AA1) were cultured according to the culture conditions described in the ATCC catalogue (http://www.atcc.org/). The cells were harvested separately, and mRNA was extracted from each cell by the method described in the literature (Molecular Cloning 2nd edition. (1989) Sambrook J., Fritsch, E.F., and Maniatis T., Cold Spring Harbor Laboratory Press). Furthermore, poly(A)*aRNA was purified from the mRNA using oligodT cellulose.

Similarly, human placenta (PLACE1, PLACE2, PLACE3, PLACE4), human ovary cancer tissue (OVARC1), tissues from human embryo at 10 weeks, which is enriched with head (HEMBA1), or body (HEMBB1), human mammary gland (MAMMA1), human thyroid gland (THYRO1), and primary cultured cells of human blood vessel endothelium (VESEN1) were used to extract mRNA by the method described in the literature (Molecular Cloning 2nd edition. (1989) Sambrook J., Fritsch, E.F., and Maniatis T., Cold Spring Harbor Laboratory Press). Furthermore, poly(A)[†]àRNA was purified from the mRNA using oligo-dT cellulose.

Each poly(A)*àRNA was used to construct a cDNA library by the oligo-capping method (Maruyama M. and Sugano S. (1994) Gene, 138: 171-174). Using the Oligo-cap linker (SEQ ID NO: 10464) and the Oligo-dT primer (SEQ ID NO: 10465), bacterial alkaline phosphatase (BAP) treatment, tobacco acid phosphatase (TAP) treatment, RNA ligation, the first strand cDNA synthesis, and removal of RNA were performed as described in the reference (Suzuki and Kanno (1996) Protein Nucleic acid and Enzyme, 41: 197-201; Suzuki Y. et al. (1997) Gene, 200: 149-156). Next, 5'- and 3'-PCR primers (SEQ ID NO: 10466, and 10467, respectively) were used for performing PCR to convert the cDNA into double stranded cDNA, which was then digested with Sfil. Then, the Dralli-cleaved pUC19FL3 vector (Figure 1; for NT2RM1, and NT2RP1), or the Drallicleaved pME18SFL3 (Figure 1) (GenBank AB009864, expression vector; for NT2RM2, NT2RM4, NT2RP2, NT2RP3, NT2RP4, NT2RP5, SKNMC1, Y79AA1, PLACE1, PLACE2, PLACE3, PLACE4, OVARC1, HEMBA1, HEMBB1, MAMMA1, THYRO1, and VESEN1) was used for cloning the cDNA in a unidirectional manner, and cDNA libraries were obtained. The nucleotide sequence of the 5'and 3'- ends of the cDNA clones was analyzed with a DNA sequencer (ABI PRISM 377, PE Biosystems) after sequencing reactions were performed with the DNA sequencing reagents (Dye Terminator Cycle Sequencing FS Ready Reaction Kit, dRhodamine Terminator Cycle Sequencing FS Ready Reaction Kit, or BigDye Terminator Cycle Sequencing FS Ready Reaction Kit, PE Biosystems), according to the instructions. The data were compiled into a database.

The full-length-enriched cDNA libraries except those for NT2RM1 and NT2RP1 were constructed using eukaryotic expression vector pME18SFL3. The vector contains $SR\alpha$ promoter and SV40 small t intron in the upstream of the cloning site, and SV40 polyA added signal sequence site in the downstream. As the cloning site of pME18SFL3 has asymmetrical Dralll sites, and the ends of cDNA fragments contain Sfil sites complementary to the Dralll sites, the cloned cDNA fragments can be inserted into the downstream of the $SR\alpha$ promoter unidirectionally. Therefore, clones containing full-length cDNA can be expressed transiently by introducing the obtained plasmid directly into COS cells. Thus, the clones can be analyzed very easily in terms of the proteins that are the gene products of the clones, or in terms of the biological activities of the proteins.

Herein, the cDNA libraries and the name of each clone are related as shown in Table 3. Therein, "xxxxxx" represents the clone number of six digits. Thus, the sequences are named by the library name, the clone number plus F- for the 5'-end, or R- for the 3'-end.

		Table 3
library:		
clone	5'-end sequence	3'-end sequence
NT2RM1:		
NT2RM1xxxxxx	F-NT2RM1xxxxxx	
NT2RP1:		
NT2RP1xxxxxx	F-NT2RP1xxxxxx	
NT2RM2:		
NT2RM2xxxxxx	F-NT2RM2xxxxxx	R-NT2RM2xxxxxx
NT2RM4:		
NT2RM4xxxxxx	F-NT2RM4xxxxxx	R-NT2RM4xxxxxx

	NT2RP2:		
	NT2RP2xxxxxx	F-NT2RP2xxxxxx	R-NT2RP2xxxxxx
_	NT2RP3:		
5	NT2RP3xxxxxx	F-NT2RP3xxxxxx	R-NT2RP3xxxxxx
	NT2RP4:		
	NT2RP4xxxxxx	F-NT2RP4xxxxxx	R-NT2RP4xxxxxx
10	NT2RP5:		
	NT2RP5xxxxxx	F-NT2RP5xxxxxx	R-NT2RP5xxxxxx
	SKNMC1:		
	SKNMC1xxxxxx	F-SKNMC1xxxxxx	R-SKNMC1xxxxxx
15	Y79AA1:		
	Y79AA1xxxxxx	F-Y79AA1xxxxxx	R-Y79AA1xxxxxx
	PLACE1:		
•	PLACE1xxxxxx	F-PLACE1xxxxxx	R-PLACE1xxxxxx
20	PLACE2:		
	PLACE2xxxxxx	F-PLACE2xxxxxx	R-PLACE2xxxxxx
	PLACE3:		
25	PLACE3xxxxxx	F-PLACE3xxxxxx	R-PLACE3xxxxxx
	PLACE4:		,
	PLACE4xxxxxx	F-PLACE4xxxxxx	R-PLACE4xxxxxx
	OVARC1:		,
30	OVARC1xxxxxx	F-0VARC1xxxxxx	R-OVARC1xxxxxx
	HEMBA1:		
	HEMBA1xxxxxx	F-HEMBA1xxxxxx	R-HEMBA1xxxxxx
35	HEMBB1:		
33	HEMBB1xxxxxx	F-HEMBB1xxxxxx	R-HEMBB1xxxxxx
	MAMMA1:		o MAINGAA
	MAMMA1xxxxxx	F-MAMMA1xxxxxx	R-MAMMA1xxxxxx
40	THYRO1:	E TIN/001	D. TUVDO1
	THYR01xxxxxx	F-THYR01xxxxxx	R-THYR01xxxxxx
	VESEN1:	r vectus	D VECENTARA
	VESEN1xxxxxx	F-VESEN1xxxxxx	R-VESEN1xxxxxx
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EXAMPLE 2

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Estimation of the fullness ratio at the 5'-ends of the clones contained in the cDNA libraries constructed by the oligo-capping method.

The fullness ratio at the 5'-end sequence of the 59,823 clones in the human cDNA libraries constructed by the oligo-capping method was determined as follows. Of all the clones whose 5'-end sequences were found in those of known human mRNA in the public database, a clone was

judged to be "full-length", if it had a longer 5'-end sequence than that of the known human mRNA, or, even though the 5'-end sequence was shorter, if it contained the translation initiation codon. A clone which did not contain the translation initiation codon was judged to be "not-full-length". The fullness ratio ((the number of full-length clones)/(the number of full-length and not-full-length clones)) at the 5'-end of the cDNA clones from each library was determined by comparing with the known human mRNA. As a result, the fullness ratio of the 5'-ends was 63.5%. The result indicates that the fullness ratio at the 5'-end sequence was extremely high.

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EXAMPLE 3

Assessment of the fullness ratio of the 5'-end of the cDNA by the ATGpr and the ESTiMateFL.

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The ATGpr, developed by Salamov A.A., Nishikawa T., and Swindells M.B. in the Helix Research Institute, is a program for prediction of the translation initiation codon based on the characteristics of the sequences in the vicinity of the ATG codon. The results are shown with expectations (also described as ATGpr1 below) that an ATG is a true initiation codon (0.05-0.94) (can be described as ATGpr1). When the program was applied to the 5'-end sequences of the clones from the cDNA library that was obtained by the oligo-capping method and that had 65% fullness ratio, the sensitivity and specificity of estimation of a full-length clone (clone containing the N-terminal end of ORF) were improved to 82-83% by selecting only clones having the ATGpr1 score 0.6 or higher.

Furthermore, the program was used to assess the fullness of 18,959 clones in the human libraries obtained here, which have 5'-ends matched to a known human mRNA.

Briefly, the maximal ATGpr1 score of the clones was determined, and then their 5'-end sequence was compared with the known human mRNA to estimate whether the clone is full-length or not. The result is shown in Table 4.

Based on the knowledge that known mRNAs, in general, are highly expressed in the cell, similar estimation was performed with genes having a low number in the EST hit, which represent relatively low abundant mRNAs, and the result is shown in Table 5.

In the table, the number of full-length clones indicate that of clones containing the N-terminal end of ORF, and so does the number of not-full-length clones that of clones without the N-terminal end of ORF. The fullness ratio represents (the number of full-length clones)/(the number of full-length clones plus the number of not-full-length clones).

Table 4

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The maximal ATGpr1 score and the fullness ratio of the 5'-end sequences of clones obtained from human cDNA libraries constructed by the oligo-capping method; clones having a matched 5'-end with that of a known mRNA.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full- length clones	fullness ratio
>=0.70	11,193	9,346	83.5%
>=0.50	13,369	10,549	78.9%
>=0.30	15,489	11,340	73.2%
>=0.15	17,394	11,811	67.9%

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The maximal ATGpr1 score and the fullness ratio of the 5'-end sequences of clones obtained from human cDNA libraries constructed by the oligo-capping method; clones having a matched 5'-end with that of a known mRNA

5'-end with that of a known mixto.			1
maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full- length clones	fullness ratio
	18.959	12.046	63.5%
>=0.00	10,939	1	

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Table 5

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequences of the clones obtained from human cDNA libraries constructed by the oligo-capping method; clones having 5 EST hits or less among the clones having a matched 5'-end with that of a known mRNA

number of full-

length clones

fullness ratio

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maximal ATGpr1

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clones plus not-full-length score clones) 69.0% 1.934 2,801 >=0.70 65.0% 2,393 3,683 >=0.50 57.8% 2,707 4,683 >=0.30 52.0% 2.890 5,559 >=0.15 49.8% 3,013 6,113 >=0.00

number of (full-length

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Next, the ESTiMateFL was used for the estimation. The ESTiMateFL, developed by Nishikawa and Ota in the Helix Research Institute, is a method for the selection of a clone with high fullness ratio by comparing with the 5'-end or 3'-end sequences of ESTs in the public database.

By the method, a cDNA clone is judged to be most likely not to be full-length if there exist any ESTs which have longer 5'-end or 3'-end sequences than the clone. The method is systematized for high throughput analysis. A clone is judged to be full-length if the clone has a longer 5'-end sequence than ESTs in the public database. Even if a clone has a shorter 5'-end, the clone is judged to be full-length if the difference in length is within 50 bases, and otherwise judged not to be full-length, for convenience.

In case of the clones whose 5'-end sequence is matching with the known mRNA, 80% of the clones judged to be full-length by comparing with ESTs was also judged to be full-length by comparing with the known mRNA. Also, 80% of the clones judged to be not full-length by comparing with ESTs was also judged to be not full-length by comparing with the known mRNA.

The precision of the estimation by comparing with ESTs is improved with increasing number of ESTs to be compared. However, in case that a limited number of ESTs are available, the reliability becomes low. Thus, the method is effective in excluding clones with high probability of being not-full-length, from the cDNA clones that is synthesized by the oligo-capping method

and that have the 5'-end sequences with about 60 % fullness ratio. In particular, the ESTiMateFL is efficiently used to estimate the fullness ratio at the 3'-end sequence of cDNA of a human unknown mRNA which has a significant number of EST deposits in the public database.

The 18,959 clones isolated from human cDNA libraries constructed by the oligo-capping method, which have the 5'-end sequence matched with a known human mRNA, were estimated by using the ATGpr and ESTiMateFL. Briefly, the 5'-end sequence of the respective clone was analyzed to obtain the maximal ATGpr1 score, and compared with the ORF of the known human mRNA that matches with it to determine whether the clone is full-length or not. Then, the 5'-end sequence of the respective clone was analyzed by the ESTiMateFL to judge whether the clone is full-length or not. Specifically, the 5'-end sequences of the 18,959 clones were compared with those of ESTs by the ESTiMateFL and the clones other than those that are not full-length were selected. Then, the selected clones were used to analyze the relationship between the ATGpr and the fullness ratio. The result was summarized in Table 6. Also, among the selected, the clones in which the number of the EST hit is not more than 5 were selected and analyzed. The result was summarized in Table 7, which represents the result of the analysis of mRNA with relatively low abundance.

Therein, the number of being full-length, the number of being not full-length, and the fullness ratio indicate the number of the clones that contain the N-terminus of the ORF, the number of the clones that do not contain the N-terminus of the ORF, and (the number of being full-length)/(the number of being full-length) plus (the number of being not full-length), respectively.

Table 6

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequence in the clones isolated from human cDNA libraries constructed by the oligo-capping method, which have the 5'-end sequence matched with a known human mRNA, and also other than those being not full-length according to the comparison with ESTs.

maximal ATGpr1 score	number of (full-length clones plus not-full- length clones)	number of full- length clones	fullness ratio
>=0.70	9,068	8,349	92.1%
>=0.50	10,345	9,318	90.1%
>=0.30	11,425	9,964	87.2%
>=0.15	12,254	10,335	84.3%
>=0.00	12,785	10,484	82.0%

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Table 7

Maximal ATGpr1 score and fullness ratio of the 5'-end sequence of the clones, which were isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA, in which the number of the EST hit is not more than 5.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full- length clones	fullness ratio
>=0.70	1,959	1,510	77.1%
>=0.50	2,469	1,821	73.8%
>=0.30	2,975	2,046	68.8%
>=0.15	3,368	2,164	64.3%
>=0.00	3,661	2,226	60.8%

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The 19,226 clones, isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to that of a known human mRNA were estimated by the ATGpr2, and the correlation between the score and the fullness ratio was estimated. Specifically, the maximal ATGpr2 score of the clones identical to a known human mRNA was determined, and then their fullness ratio was estimated by comparing the 5'-ends with ORF of known human mRNA. The result was shown in Table 8.

Table 8

Maximal ATGpr2 score and fullness ratio of the 5'-end sequence of the clones, which are isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA.

maximal ATGpr2 score	number of (full-length clones plus not-full-length clones)	number of full- length clones	fullness ratio
>=0.30	10,748	8,031	74.7%
>=0.15	16,383	11,226	68.5%
>=0.00	19,226	12,285	63.9%

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According to the above results, it was found that, in case of using clones isolated from human cDNA libraries constructed by the oligo-capping method, the fullness ratio of the clones that have low score in the ATGpr can be improved by estimating their 5'-end sequence using the combination of the ATGpr and the ESTiMateFL. Therefore, the method was applied to select a cDNA clone with high fullness ratio.

EXAMPLE 4

Clustering of the 5'-end and 3'-end sequences of cDNA clones.

The 5'-end and 3'-end sequences of cDNA clones were obtained, and clustered separately. The single pass data of the nucleotide sequence of the 5'-end and 3'-end was subject to the BLAST search between the sequence data of all the clones synthesized in example 1, and the clones considered to be originating from the same gene were put together into a group. If the 5'-end of a clone contains the consensus sequence of 300 bases or more with identity 95% or more, or the 3'-end contains the consensus sequence of 200 bases or more and having identity 90% or more, the clones were put in the same group.

The groups of the 5'-end sequence and the 3'-end sequence were further clustered so as that the groups from the same clone can be in the same group (cluster).

EXAMPLE 5

20 Characterization of the cloned sequence.

The data of the 5'-end sequence of the cloned sequence was characterized by the following way:

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(1) examining whether it is identical to those of mRNA from human and other species (including authorized sequences) and human EST by the BLAST homology search of the GenBank,

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(2) examining whether it has longer 5'-end than those of human mRNA and human EST.

(3) determining the scores in the ATGpr1 and ATGpr2 programs of all the initiation codons in the 5'-end sequence, and

(4) determining the number of the human EST clone(s) that is judged to be identical by the BLAST homology search of the GenBank.

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- The data of the 3'-end sequence of the cloned sequence was characterized by the above (1) and (4).
- These characterized data were used for the final selection of the clones.

EXAMPLE 6

ldentity to the human mRNA and human EST, and comparison of the length of the 5'-end.

The 5'-end and 3'-end sequences of the cloned sequence was judged to be identical to those of mRNA from human or other species when the sequence to compare has the length of 200 bases or longer, and the obtained homology is 94% or more. The 5'-end and 3'-end sequences of the cloned sequence was judged to be identical to those of human EST when the sequence to

compare has the length of 200 bases or longer, and the obtained homology is 90% or more.

The sequence of the clone was judged to be full-length in comparison with human mRNA when the sequence has longer 5'-end, or it contains the translation initiation site. The sequence of the clone was judged to be full-length in comparison with human EST in the database when the sequence has longer 5'-end, or while it has shorter end, the difference in length between the two sequences is 50 bases or less. The other clones were judged to be not full-length.

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EXAMPLE 7

Prediction of the fullness ratio by the ATGpr.

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The score in the ATGpr1 is the expectation to be full-length based on calculations, and the higher score reflects the higher probability to be full-length as shown in Example 3. The maximal ATGpr1 score and the maximal ATGpr2 score represent the score obtained with all the initiation codons contained in the 5'-end sequence of the cloned sequence, and were used for the characterization.

EXAMPLE 8

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Prediction of the novelty using the number of the identical ESTs by the homology search.

For both the 5'-end and 3'-end sequences of the clones, the number of the identical ESTs was determined by the homology search on the GenBank. Human ESTs were judged to be identical when the EST has a sequence of 200 nucleotides or more with 90% or more matching with the 5'-end sequence. The number of the identical ESTs were used for characterization and as an index of novelty. The clone having not identical sequence at the 5'-end and 3'-end sequences to those of mRNA as well as those of ESTs is a gene encoding a novel protein. Similarly, a clone having either the 5'-end or the 3'-end sequences, which has low number of the identical ESTs, is judged to be a gene encoding a novel protein.

40 EXAMPLE 9

Characterization of clusters.

sequence.

- The clusters of the groups of the 5'-end and 3'-end sequences were characterized according to the following criteria.
 - (1) Whether it is identical to the mRNA sequences from human or other species (including authorized sequences), or human ESTs by the BLAST search of the GenBank.
 A cluster containing at least one sequence of all the 5'-end and 3'-end sequences, which is identical to one of the mRNA sequences, was regarded to be the same cluster of the mRNA

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(2) Whether it has longer 5'-end than human mRNA sequence and human ESTs. When all the 5'-end sequences contained in a cluster are judged to be not full-length

compared with the mRNA sequences and human ESTs, the cluster was regarded as being not full-length.

5 (3) The scores in the ATGpr1 and ATGpr2 using all the initiation codons contained in the 5'end sequences.

The maximal ATGpr1 score among those of all the 5'-end sequences in a cluster was determined as the ATGpr1 score of the cluster. The ATGpr2 score of the cluster was also determined in the same way.

(4) The number of the identical human ESTs determined by the BLAST search of the GenBank

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The maximum number was determined in the numbers of ESTs identical to each of 5'-end sequences contained in a cluster. The number of the ESTs identical to the 5'-end sequences in the cluster was defined as the maximum number. The number of the ESTs identical to the 3'-end sequences in a cluster was determined in the same way.

EXAMPLE 10

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Methods for selection of the clusters by the characteristics.

Data obtained by the characterization described above was used to discard the clusters that are identical to any mRNA sequence from human and other species (including authorized sequences), or those clusters that are not full-length. From the rest of the clusters, the clusters that fulfill any of the following conditions were selected.

- (a) A cluster in which the number of the identical ESTs for the 5'-end sequence is 20 or less, and the ATGpr1 score exceeds 0.3.
- (b) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for both the 5'-end sequence and the 3'-end sequence is 5 or less, and multiple clones are contained.
- (c) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for the 5'-end sequence is 0, and the number of the identical ESTs for the 3'-end sequence is not less than 1.
- (d) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for the 5'-end sequence is not less than 1 and not more than 5, and the number of the identical ESTs for the 3'-end sequence is 0.

The clusters selected by (a) contain at least one clone that is novel and having high fullness ratio. The clusters selected by (b), (c), and (d) contain at least one clone that is novel and having low fullness ratio, but is still full-length.

EXAMPLE 11

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5 Methods for selection of clones from clusters.

In the clusters comprising a single clone, the clone was selected.

In the clusters comprising multiple clones, in which multiple clones have the ATGpr1 score higher than 0.3, a clone with the highest score was selected.

In the clusters comprising multiple clones, in which multiple clones have the ATGpr1 score not more than 0.3, a clone with the highest ATGpr2 score was selected, if the score was higher than 0.3.

In the clusters comprising multiple clones, in which the clones have the scores not more than 0.3 in both the ATGpr1 and the ATGpr2, a clone with the highest scores in both the ATGpr1 and ATGpr2 was selected.

In the clusters comprising multiple clones, in which the above selection by the ATGpr score was not applicable, selected was a clone having longer 5'-end by assembling the 5'-end sequence, 3'-end sequence, and human ESTs. For assembling, the Sequencher™ (Hitachi Soft Engineering) was used. When even the selection by assembling failed, all the clones were judged to be full-length.

As a result, 3690 clones were the clones that have the maximal ATGpr1 score higher than 0.3. On the other hand, 477 clones were the clones that have the maximal ATGpr1 score not more than 0.3, and the maximal ATGpr2 score higher than 0.3. The number of the clones having the highest scores in both the ATGpr1 and ATGpr2, while the scores were not more than 0.3, were 97. The number of the clones which were not selected by the ATGpr scores, but were selected by assembling the 5'-end sequence, 3'-end sequence, and human ESTs, were 117. The clones that have the score in both the ATGpr1 and ATGpr2 not more than 0.3, but were selected because the cluster comprises a single clone, were 1166. In the clones, at least either of the 5'-end or 3'end sequence was not identical to any of human ESTs. Some clones were selected because the cluster comprises a single clone, or by assembling, in which there is no ATG codon (9 clones: HEMBA106569, HEMBB1001454, NT2PR2002839, NT2RP2005325, HEMBA1001960 NT2RP2006323, PLACE1004506, PLACE1005526, and THYRO1001177). The sequences that do not contain the ATG codon were considered to be corresponding to the 5'-UTR. Although the clones do not have the scores in the ATGpr1 and ATGpr2, the clones were yet judged to be fulllength according to the fullness ratio, as shown in Table 4, 5, 6, 7, and 8. The above clones that were finally judged to be full-length were classified into 11 groups according to the following criteria.

Group (1): 1516 clones

Among the 3690 clones having the maximal ATGpr1 score higher than 0.3, the following 1516 clones were having high fullness ratio and a novel clone, in which at least either of the 5'-end or 3'-end sequence, or both of them were not identical to any of human ESTs.

HEMBA1000046, HEMBA1000050, HEMBA1000129, HEMBA1000150, HEMBA1000158, HEMBA1000193 , HEMBA1000201, HEMBA1000216, HEMBA1000227, HEMBA1000288, HEMBA1000290, HEMBA100030 3, HEMBA1000304, HEMBA1000369, HEMBA1000392, HEMBA1000396, HEMBA1000488, HEMBA10005 05, HEMBA1000508, HEMBA1000534,

HEMBA1000542, HEMBA1000594, HEMBA1000637, HEMBA1000 657, HEMBA1000752, HEMBA1000867. HEMBA1000869. HEMBA1000872. HEMBA1000910. HEMBA100 0918. HEMBA1000919, HEMBA1000942, HEMBA1000968, HEMBA1000975, HEMBA1000986, HEMBA10 01022, HEMBA1001043, HEMBA1001052, HEMBA1001080, HEMBA1001085, HEMBA1001109, HEMBA1 001140, HEMBA1001174, HEMBA1001235, HEMBA1001286, HEMBA1001302, HEMBA1001398, HEMBA 1001407, HEMBA1001415, HEMBA1001446, HEMBA1001476, HEMBA1001497, HEMBA1001510, HEMB A1001533, HEMBA1001570, HEMBA1001581, HEMBA1001635, HEMBA1001640, HEMBA1001647, HEM BA1001661, HEMBA1001731, HEMBA1001744, HEMBA1001746, HEMBA1001800, HEMBA1001815, HE MBA1001822, HEMBA1001866, HEMBA1001896, HEMBA1001910, HEMBA1001987, HEMBA1002018. HEMBA1002035. HEMBA1002049. HEMBA1002092. HEMBA1002119. HEMBA1002125 , HEMBA1002161, HEMBA1002177, HEMBA1002189, HEMBA1002191, HEMBA1002199, HEMBA100222 9, HEMBA1002237, HEMBA1002265, HEMBA1002363, HEMBA1002417, HEMBA1002419, HEMBA10024 30, HEMBA1002439, HEMBA1002477, HEMBA1002503, HEMBA1002508, HEMBA1002515, HEMBA1002 547, HEMBA1002688, 20 HEMBA1002703, HEMBA1002746, HEMBA1002750, HEMBA1002850, HEMBA100 2973, HEMBA1003021, HEMBA1003067, HEMBA1003077, HEMBA1003078, HEMBA1003079, HEMBA10 03117, HEMBA1003129, HEMBA1003175, HEMBA1003199, HEMBA1003235, HEMBA1003250, HEMBA1 003257, HEMBA1003291, HEMBA1003322, HEMBA1003327, HEMBA1003370, HEMBA1003380, HEMBA 1003395, HEMBA1003402, HEMBA1003461, HEMBA1003480, HEMBA1003538, HEMBA1003545, HEMB A1003556, HEMBA1003581, HEMBA1003621, HEMBA1003645, HEMBA1003667, HEMBA1003720, HEM BA1003760, HEMBA1003799, HEMBA1003807, HEMBA1003827, HEMBA1003836, HEMBA1003866, HE MBA1003879, HEMBA1003880, HEMBA1003985, HEMBA1003989, HEMBA1004011, HEMBA1004048, HEMBA1004056, HEMBA1004074, HEMBA1004133, HEMBA1004146 , HEMBA1004150, HEMBA1004199, HEMBA1004200, HEMBA1004238, HEMBA1004246, HEMBA100427 5, HEMBA1004286, HEMBA1004289, HEMBA1004321, HEMBA1004327, HEMBA1004335, HEMBA10043 41, HEMBA1004372, HEMBA1004389, HEMBA1004479, HEMBA1004499, HEMBA1004507, HEMBA1004 542, HEMBA1004554, HEMBA1004573, HEMBA1004596, HEMBA1004632, HEMBA1004697, HEMBA100 4705, HEMBA1004709, HEMBA1004711, HEMBA1004736, HEMBA1004751, HEMBA1004752, HEMBA10 04753, HEMBA1004756, HEMBA1004758, HEMBA1004768, HEMBA1004771, HEMBA1004795, HEMBA1 004806, HEMBA1004850, HEMBA1004863, HEMBA1004889, HEMBA1004923, HEMBA1004929, HEMBA 1004930, HEMBA1004944, HEMBA1004972, HEMBA1004980, HEMBA1005019, HEMBA1005035, HEMB A1005050, HEMBA1005066, HEMBA1005075, HEMBA1005079, HEMBA1005083, HEMBA1005113, HEM BA1005133, HEMBA1005149, HEMBA1005219, HEMBA1005331, HEMBA1005338, HEMBA1005367, HE HEMBA1005468, HEMBA1005469, HEMBA1005517, HEMBA1005518, MBA1005411, HEMBA1005526, H EMBA1005548, HEMBA1005576, HEMBA1005583, HEMBA1005595, HEMBA1005609, HEMBA1005685, HEMBA1005732, HEMBA1005755, HEMBA1005813, HEMBA1005815, HEMBA1005834, HEMBA1005884, HEMBA1005963, HEMBA1005991, HEMBA1006031, HEMBA1006100, HEMBA1006121, HEMBA1006138, HEMBA1006173, HEMBA1006182 , HEMBA1006198, HEMBA1006252, HEMBA1006272, HEMBA1006278, HEMBA1006291, HEMBA100629 3, HEMBA1006344, HEMBA1006349, HEMBA1006377,

HEMBA1006381, HEMBA1006398, HEMBA10064 24, HEMBA1006467, HEMBA1006474, HEMBA1006483, HEMBA1006492, HEMBA1006494, HEMBA1006 497, HEMBA1006502,

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HEMBA1006530, HEMBA1006579, HEMBA1006583, HEMBA1006643, HEMBA100 6674,
      HEMBA1006682, HEMBA1006709, HEMBA1006717, HEMBA1006737, HEMBA1006754,
      HEMBA10 06758, HEMBA1006767, HEMBA1006795, HEMBA1006796,
                                                                                                     HEMBA1006807,
      HEMBA1006832, HEMBA1 006900, HEMBA1006973, HEMBA1006976, HEMBA1006993,
      HEMBA1006996, HEMBA1007002, HEMBA 1007052, HEMBA1007062,
                                                                                                     HEMBA1007066,
      HEMBA1007151, HEMBA1007203, HEMBA1007281, HEMB A1007300,
                                                                                                     HEMBA1007320,
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      HEMBB1000008,
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- PLACE10 09404, PLACE1009434, PLACE1009443, PLACE1009459, PLACE1009524, PLACE1009542, PLACE1 009607, PLACE1009798, PLACE1009861, PLACE1009971, PLACE1009992, PLACE1009997, PLACE 1010148, PLACE1010261, PLACE1010293, PLACE1010310, PLACE1010321, PLACE1010324, PLAC E1010329, PLACE1010401,
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 PLACE2000047, PLACE2000050, PLACE2000062, PLACE2000072,

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	PLACE2000223,	PLACE2000235, PLACE20002 46, PLACE2000274, PLACE2000359,
	PLACE2000371,	PLACE2000373, PLACE2000411, PLACE2000 433, PLACE2000438,
5	PLACE2000458,	
	PLACE3000020,	PLACE3000029, PLACE3000059, PLACE3000070, PLACE3000103,
	PLACE3000142	, PLACE3000145, PLACE3000147, PLACE3000155, PLACE3000157,
	PLACE3000194,	PLACE300022 1, PLACE3000244, PLACE3000254, PLACE3000304,
10	PLACE3000310,	PLACE3000320, PLACE30003 22, PLACE3000331, PLACE3000350,
	PLACE3000352,	PLACE3000363, PLACE3000405, PLACE3000 416, PLACE3000425,
	PLACE3000455,	· · · · · · · · · · · · · · · · · · ·
	PLACE4000014,	PLACE4000052, PLACE4000063, PLACE4000128, PLACE4000129,
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	PLACE4000222,	PLACE400023 0, PLACE4000261, PLACE4000270, PLACE4000323,
	PLACE4000387,	PLACE4000392, PLACE40004 50, PLACE4000465, PLACE4000487,
	PLACE4000489,	PLACE4000521, PLACE4000522, PLACE4000 590, PLACE4000612,
20	PLACE4000638,	PLACE4000654,
	SKNMC1000011,	SKNMC1000013, THYRO1000034, THYRO1000070, THYRO1000072.
	THYRO1000107	, THYRO1000124, THYRO1000163, THYRO1000253, THYRO1000343,
	THYRO1000381,	THYRO100038 7, THYRO1000395, THYRO1000501, THYRO1000505,
25	THYRO1000637,	THYRO1000666, THYRO10006 76, THYRO1000684, THYRO1000734,
	THYRO1000787,	THYRO1000916, THYRO1000934, THYRO1000 951, THYRO1001033,
	THYRO1001100,	THYRO1001142, THYRO1001189, THYRO1001213, THYRO100 1320,
	THYRO1001374,	THYRO1001403, THYRO1001458, THYRO1001487, THYRO1001602,
30	THYRO10 0160	5, THYRO1001637, THYRO1001656, THYRO1001738, THYRO1001809,
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	Y79AA1000627,	Y79AA1000824, Y79AA 1000827, Y79AA1000850, Y79AA1000962,
	Y79AA1000976,	Y79AA1001068, Y79AA1001145, Y79A A1001185, Y79AA1001233,
35	Y79AA1001299,	Y79AA1001391, Y79AA1001541, Y79AA1001555, Y79 AA1001581,
	Y79AA1001613,	Y79AA1001846, Y79AA1001866, Y79AA1001874, Y79AA1002115, Y7
	9AA1002139, Y79	AA1002229, Y79AA1002246, Y79AA1002298, Y79AA1002361, Y79AA1002399,
		Y79AA1002433, Y79AA1002472, Y79AA1002482,
40	·	·

Group (2): 377 clones

Among the 3690 clones, the following 377 clones were full-length, and a novel clone, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

HEMBA1000156, HEMBA1000231, HEMBA1000244, HEMBA1000302, HEMBA1000523, HEMBA1000531 , HEMBA1000555, HEMBA1000568, HEMBA1000971, HEMBA1001009, HEMBA1001133, HEMBA100113 7, HEMBA1001257, HEMBA1001281, HEMBA1001289, HEMBA1001310, HEMBA1001377, HEMBA10013 88, HEMBA1001413, HEMBA1001455, HEMBA1001579, HEMBA1001702, HEMBA1001781, HEMBA1001 804, HEMBA1001819, HEMBA1001864, HEMBA1001869, HEMBA1001912, HEMBA1001915, HEMBA100 2084, HEMBA1002102, HEMBA1002212, HEMBA1002495, HEMBA1002513, HEMBA1002863, HEMBA10 02935, HEMBA1002970, HEMBA1002971, HEMBA1002997, HEMBA1003680, HEMBA1003692, HEMBA1003804, HEMBA 1003893, HEMBA1003953, HEMBA1004097,

- HEMBA1004207, HEMBA1004502, HEMBA1004560, HEMB A1004638, HEMBA1004693, HEMBA1004973, HEMBA1004977, HEMBA1005223, HEMBA1005244, HEM BA1005314, HEMBA1005680, HEMBA1005737, HEMBA1005746, HEMBA1005822, HEMBA1005891, HEMBA1005
- 5 MBA1006445, HEMBA1006535, HEMBA1006566, HEMBA1006877, HEMBA1006936, HEMBA1007178, H EMBA1007251, HEMBA1007267,
 - HEMBB1000030, HEMBB1000215, HEMBB1000226, HEMBB1000449, HEMBB1000789, HEMBB1001153, HEMBB1001175, HEMBB1001234, HEMBB1001387,
- HEMBB1001905, HEMBB100190 6, HEMBB1002300, HEMBB1002383, HEMBB1002489,
 HEMBB1002607, HEMBB1002705, MAMMA1000257, MAMMA1000416, MAMMA1000831,
 MAMMA1000841, MAMMA1001021, MAMMA1001041, MAMMA1001105, MAMMA1001139,
 MAMMA1001198, MAMMA1001730, MAMMA1001837, MAMMA100219 8, MAMMA1002209.
- MAMMA1002219, MAMMA1002292, MAMMA1002297, MAMMA1002637, MAMMA10026 50,
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- NT2RM4001032, NT2RM400104 7, NT2RM4001116, NT2RM4001140, NT2RM4001200, NT2RM4001384, NT2RM4001412, NT2RM40014 83, NT2RM4001715, NT2RM4001783, NT2RM4001979, NT2RM4002034, NT2RM4002044, NT2RM4002 145, NT2RM4002344, NT2RM4002452,
- 25 NT2RP2000006, NT2RP2000008, NT2RP2000054, NT2RP2000079, NT2RP2000097. NT2RP2000126 NT2RP2000224, NT2RP2000274, NT2RP2000764, NT2RP2000812, NT2RP2000819. NT2RP200084 5. NT2RP2000863. NT2RP2000880, NT2RP2001127, NT2RP2001218 NT2RP2001233, NT2RP20013 94. NT2RP2001420, NT2RP2001576.
- NT2RP2001597, NT2RP2001721, NT2RP2001907, NT2RP2002 046, NT2RP2002312, NT2RP2002325, NT2RP2002503, NT2RP2002537, NT2RP2002862, NT2RP200 2925, NT2RP2002939, NT2RP2002954, NT2RP2002986, NT2RP2003034, NT2RP2003125, NT2RP20 03286, NT2RP2003293, NT2RP2003347, NT2RP2003433, NT2RP2003445,
- 35 NT2RP2003543, NT2RP2 003781, NT2RP2004066, NT2RP2004316, NT2RP2004392, NT2RP2004897, NT2RP2004999, NT2RP 2005003, NT2RP2005393. NT2RP2005520, NT2RP2005752, NT2RP2005722, NT2RP2005763, NT2R P2006023, NT2RP2006186, NT2RP2006565,
- NT2RP3000590, NT2RP3000527, NT2RP3000562, NT2RP3000596, NT2RP3000599, NT2RP3001004 NT2RP3001126. NT2RP3001150. NT2RP3001392. NT2RP3001407. NT2RP3001457, NT2RP300175 NT2RP3001929, 3. NT2RP3001969, NT2RP3002151, NT2RP3002303. NT2RP3002484. NT2RP30025 90. NT2RP3002660. NT2RP3003078.
- NT2RP3003204, NT2RP3003251, NT2RP3003411, NT2RP3003 831, NT2RP3003870, NT2RP3004253, NT2RP3003876, NT2RP3004093. NT2RP3004246, NT2RP300 4348, NT2RP3004504, NT2RP3004507, NT2RP4000312, NT2RP4000518, NT2RP4000781, NT2RP40 00975, NT2RP4000997, NT2RP4002408, NT2RP5003512,
- OVARC1000060, OVARC1000411, OVARC1000440, OVARC1000461, OVARC1000703, OVARC1000846, OVARC1000876, OVARC1000915, OVARC1001040, OVARC1001113, OVARC1001167, OVARC100117 0, OVARC1001188, OVARC1001271, OVARC1001329, OVARC1001610, OVARC1001767, OVARC10018 20, OVARC1001879, OVARC1002138,
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- PLACE1001570, PLACE1001602, PLACE10016 10, PLACE1001632, PLACE1001989. PLACE1002115. PLACE1002465, PLACE1002532. PLACE1002 851, PLACE1002991, PLACE1002993, PLACE1003025, PLACE1003342, PLACE1003553, PLACE100 3605, PLACE1003669, PLACE1003723, PLACE1003738, PLACE1003762, PLACE1003783,
- PLACE10 04104, PLACE1004114, PLACE1004149, PLACE1004197, PLACE1004473, PLACE1004510, PLACE1 004645, PLACE1004672, PLACE1004868,
- PLACE1005374, PLACE1005313, PLACE1005834, PLACE1005921, PLACE1005936, PLACE1006239, PLACE1005966 PLACE1006157, PLACE1006225, PLACE1006248, PLACE100653 1, PLACE1006626, PLACE1006445, PLACE1006731, PLACE1006901, PLACE1006961, PLACE1006966, PLACE10073 17, PLACE1007375, PLACE1007460, PLACE1007544, PLACE1007583, PLACE1007791, PLACE1007 843, PLACE1007897,
- PLACE1008129, PLACE1008309, PLACE1008401, PLACE1008402, PLACE100 8532, PLACE1008627, PLACE1008808, PLACE1008902, PLACE1008934, PLACE1009099, PLACE10 09130, PLACE1009166, PLACE1009659, PLACE1009665, PLACE1009708, PLACE10109704, PLACE1 009886, PLACE10109908, PLACE1010202, PLACE1010580, PLACE1010661,
- PLACE1010811, PLACE 1010917, PLACE1010942, PLACE1010965, PLACE1011026, PLACE1011325, PLACE1011472, PLAC E1011563, PLACE1011664, PLACE1011682, PLACE1011982, PLACE4000049, PLACE4000300, PLA CE4000558,
- THYRO1000368, THYRO1000662, THYRO1000712, THYRO1000783, THYRO1000952, 25 THYRO1000988 THYRO1001661, THYRO1001854, Y79AA1000037, Y79AA1000059, Y79AA100110, 5, Y79AA1001177, Y79AA1000985, Y79AA1001211, Y79AA1001281, Y79AA1001533, Y79AA1001923, Y79AA10020 83, Y79AA1002204, Y79AA1002407,

30 Group (3): 1797 clones

Among the 3690 clones, the following 1797 clones were full-length, and novel clones, in which the number of the identical human ESTs for the 5'-end sequence is not more than 20 (except clones in which at least either of the 5'-end or 3'-end sequence, or both of them are not

- identical to any of human ESTs, and clones in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5).
 - HEMBA100005, HEMBA1000012, HEMBA1000020, HEMBA1000030, HEMBA1000076, HEMBA1000141, HEMBA1000168, HEMBA1000185, HEMBA1000307, HEMBA1000327,
- 40 HEMBA1000356, HEMBA100038 7, HEMBA1000456, HEMBA1000460, HEMBA1000490, HEMBA1000491, HEMBA1000501, HEMBA10005 20, HEMBA1000561, HEMBA1000588, HEMBA1000591, HEMBA1000592, HEMBA1000608, HEMBA1000 636, HEMBA1000682,
- HEMBA1000686, HEMBA1000719, HEMBA1000727, HEMBA1000817, HEMBA100 0851,

 HEMBA1000946, HEMBA1001008, HEMBA1001059, HEMBA1001088, HEMBA1001122,

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- 50 HEMBA1001672, HEMBA1001711, HEMBA1001714, HEMB A1001723, HEMBA1001809, HEMBA1001824, HEMBA1001847, HEMBA1001913, HEMBA1001921, HEM BA1001939, HEMBA1001950, HEMBA1001967,
- HEMBA1002150, HEMBA1002151, HEMBA1002215, HEMBA1002241, HEMBA1002267, HEMBA1002341, HEMBA1002458, HEMBA1002460, HEMBA1002462, HEMBA1002469, HEMBA1002475, HEMBA100254 2, HEMBA1002569, HEMBA1002583, HEMBA1002609, HEMBA1002624, HEMBA1002696, HEMBA10027 68, HEMBA1002770, HEMBA1002777,

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- NT2RM2001903, NT2RM2 001935, NT2RM2001936, NT2RM2001950, NT2RM2001997, NT2RM2001998, NT2RM2002004, NT2RM 2002030, NT2RM2002049, NT2RM2002055, NT2RM2002088, NT2RM2002091, NT2RM2002100, NT2R M2002109, NT2RM2002128,

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 PLACE1006819, PLACE1006883, PLACE10069 17, PLACE1006989, PLACE1007021,

 PLACE1007053, PLACE1007105, PLACE1007178, PLACE1007 238, PLACE1007239,
- PLACE1007243, PLACE1007282, PLACE1007367, PLACE1007386, PLACE100 7402, PLACE1007409, PLACE1007416, PLACE1007454, PLACE1007507, PLACE1007511, PLACE10 07547, PLACE1007598, PLACE1007621, PLACE1007632, PLACE1007645, PLACE1007649, PLACE1 007688, PLACE1007690, PLACE1007697, PLACE1007705, PLACE1007706,
- PLACE1007725, PLACE 1007730, PLACE1007746, PLACE1007846, PLACE1007858, PLACE1007908, PLACE1007954, PLAC E1007955, PLACE1007958, PLACE1007969, PLACE1007990,
- PLACE1008044, PLACE1008122, PLACE1008132, PLACE1008198, PLACE1008209, PLACE1008356 , PLACE1008368, PLACE1008398, PLACE1008429, PLACE1008524, PLACE1008531, PLACE100856 8, PLACE1008603, PLACE1008629, PLACE1009650, PLACE1009027, PLACE1009045, PLACE10090 60, PLACE1009091, PLACE1009186, PLACE1009298, PLACE1009319, PLACE1009338, PLACE1009 444, PLACE1009468,
- 40 PLACE1009476, PLACE1009571, PLACE1009581, PLACE1009596, PLACE100 9622, PLACE1009670, PLACE1009721, PLACE1009731, PLACE1009763, PLACE1009845, PLACE10 09921, PLACE1009995,
- PLACE1010023, PLACE1010031, PLACE1010053, PLACE1010074, PLACE1010076,

 PLACE1010096 , PLACE1010102, PLACE1010105, PLACE1010106, PLACE1010134,

 PLACE1010152, PLACE101019 4, PLACE1010274, PLACE1010362, PLACE1010364,

 PLACE1010383, PLACE1010491, PLACE10104 92, PLACE1010522, PLACE1010529,

 PLACE1010599, PLACE1010622, PLACE1010630, PLACE1010 720, PLACE1010761,
- 50 PLACE1010771, PLACE1010786, PLACE1010800, PLACE1010833, PLACE101 0856, PLACE1010857, PLACE1010870, PLACE1010877, PLACE1010925, PLACE1010944, PLACE10 10960, PLACE1011041, PLACE1011090, PLACE1011160, PLACE1011214, PLACE1011219, PLACE1 011221, PLACE1011263, PLACE1011291, PLACE1011310, PLACE1011332,
- PLACE1011371, PLACE 1011433, PLACE1011477, PLACE1011586, PLACE1011646, PLACE1011675, PLACE1011858, PLAC E1011923,
 - PLACE2000007, PLACE2000021, PLACE2000030, PLACE2000164, PLACE2000172,

- PLACE2000302 , PLACE2000341, PLACE2000399, PLACE2000404, PLACE2000427, PLACE300009, PLACE300012 1, PLACE3000148, PLACE3000156, PLACE3000160, PLACE3000197, PLACE3000226, PLACE30002 42, PLACE3000339, PLACE3000353,
- PLACE3000413, PLACE3000477, PLACE4000009, PLACE4000 034, PLACE4000089, PLACE4000106, PLACE4000259, PLACE4000269, PLACE4000326, PLACE400 0369, PLACE4000431, PLACE4000445, PLACE4000581, PLACE4000593, PLACE4000650, PLACE40 00670.
- 10 SKNMC1000046, SKNMC1000050, SKNMC1000091, THYRO1000017, THYRO1000040, THYRO1000121 , THYRO1000173, THYRO1000197, THYRO1000199, THYRO1000206, THYRO1000242, THYRO100027 0, THYRO1000288, THYRO1000320, THYRO1000327, THYRO1000358, THYRO1000394, THYRO10004 01, THYRO1000488, THYRO1000502,
- THYRO1000569, THYRO1000570, THYRO1000585, THYRO1000 605, THYRO1000715, THYRO1000756, THYRO1000777, THYRO1000829, THYRO1000855, THYRO100 0926, THYRO1000983, THYRO1000984, THYRO1001120, THYRO1001134, THYRO1001173, THYRO10 01204, THYRO1001271, THYRO1001287, THYRO1001313, THYRO1001347,
- 20 THYRO1001363, THYRO1 001405, THYRO1001406, THYRO1001537, THYRO1001584, THYRO1001671, THYRO1001703, THYRO 1001721, THYRO1001793,
 - VESEN1000122, Y79AA1000013, Y79AA1000181, Y79AA1000202, Y79AA1000214, Y79AA1000230 , Y79AA1000231, Y79AA1000258, Y79AA1000313, Y79AA1000328,
- Y79AA1000342, Y79AA100036 8, Y79AA1000420, Y79AA1000469, Y79AA1000480, Y79AA1000539, Y79AA1000540, Y79AA10005 Y79AA1000574, Y79AA1000589, 60 Y79AA1000705, Y79AA1000734, Y79AA1000748, Y79AA1000 Y79AA1000774, Y79AA1000782. Y79AA1000784, Y79AA1000800, Y79AA100 Y79AA1000794, 0833.
- Y79AA1000966, Y79AA1000968, Y79AA1001023, Y79AA1001041, Y79AA1001048, Y79AA1001077, Y79AA1001078, Y79AA1001216, Y79AA1001228, Y79AA1001236, Y79AA1001312, Y79AA1 001323, Y79AA1001394, Y79AA1001402, Y79AA1001493, Y79AA1001511, Y79AA1001548, Y79AA 1001585, Y79AA1001603, Y79AA1001647, Y79AA1001665,
- Y79AA1001679, Y79AA1001696, A1001705, Y79AA1001711, Y79A Y79AA1001781, Y79AA1001805, Y79 Y79AA1001827, Y79AA1001963, AA1002027, Y79AA1002089, Y79AA1002093, Y79AA1002125, Y79AA1002208, Y79AA1002209, Y7 9AA1002210, Y79AA1002234, Y79AA1002258, Y79AA1002307, Y79AA1002311, Y79AA1002351, Y
- ⁴⁰ 79AA1002416, Y79AA1002487,

Group (4): 453 clones

- Among the 1857 clones having the maximal ATGpr1 score not more than 0.3 (including the 9 clones whose 5'-end sequence does not contain the ATG codon), the following 453 clones were judged to be full-length since their ATGpr2 score was 0.3 or higher (Table 11). The clones were novel clones, in which at least either of the 5'-end or 3'-end sequence is not identical to any of human ESTs.
- HEMBA1000180, HEMBA1000280, HEMBA1000282, HEMBA1000333, HEMBA1000351,
 HEMBA1000357 , HEMBA1000376, HEMBA1000469, HEMBA1000504, HEMBA1000519,
 HEMBA1000545, HEMBA100055 7, HEMBA1000575, HEMBA1000726, HEMBA1000822,
 HEMBA1000934, HEMBA1000943, HEMBA10009 60, HEMBA1000972, HEMBA1001020,
- HEMBA1001051, HEMBA1001060, HEMBA1001071, HEMBA1001 208, HEMBA1001319,
 HEMBA1001383, HEMBA1001411, HEMBA1001433, HEMBA1001435, HEMBA100 1478,
 HEMBA1001522, HEMBA1001636, HEMBA1001651, HEMBA1001658, HEMBA1001709,

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HEMBA10 01745, HEMBA1001791, HEMBA1001835,
                                             HEMBA1001844, HEMBA1001888,
HEMBA1001940, HEMBA1 001962, HEMBA1002022,
                                             HEMBA1002185,
                                                           HEMBA1002348,
HEMBA1002621, HEMBA1002645, HEMBA 1002661,
                                             HEMBA1002728,
                                                           HEMBA1002921,
HEMBA1002924,
              HEMBA1002934,
                             HEMBA1002968,
                                            HEMBA1003037,
                                                           HEMBA1003071,
HEMBA1003166,
              HEMBA1003202,
                                            HEMBA1003220,
                             HEMBA1003204,
                                                           HEMBA1003229,
HEMBA1003276,
              HEMBA1003296, HEMBA1003373, HEMBA1003531, HEMBA100364 0,
HEMBA1003856,
              HEMBA1003926,
                             HEMBA1003942.
                                            HEMBA1003987,
                                                           HEMBA1004024.
                                             HEMBA1004577,
HEMBA10042 67, HEMBA1004323, HEMBA1004433,
                                                           HEMBA1004730,
HEMBA1004778, HEMBA1004 803, HEMBA1004807, HEMBA1004880,
                                                           HEMBA1004900.
HEMBA1004983,
              HEMBA1005123, HEMBA100 5241, HEMBA1005311,
                                                           HEMBA1005318,
HEMBA1005353,
              HEMBA1005374, HEMBA1005447, HEMBA10 05588,
                                                           HEMBA1005593,
HEMBA1005606,
             HEMBA1005679, HEMBA1005894, HEMBA1005911, HEMBA1 006036,
HEMBA1006124, HEMBA1006253, HEMBA1006259, HEMBA1006364, HEMBA1006380, HEMBA
1006426,
          HEMBA1006562,
                          HEMBA1006597,
                                           HEMBA1006639,
                                                           HEMBA1006653,
HEMBA1006696, HEMB A1006744, HEMBA1006824, HEMBA1006949,
                                                           HEMBA1007078,
HEMBA1007129, HEMBA1007147, HEM BA1007206, HEMBA1007279, HEMBA1007327,
              HEMBB1000055, HEMBB1000144,
                                            HEMBB1000258,
HEMBB1000005.
                                                           HEMBB1000318.
HEMBB1000335,
              HEMBB1000354,
                             HEMBB1000374,
                                            HEMBB1000402,
                                                           HEMBB1000404.
HEMBB1000480,
              HEMBB100049 3, HEMBB1000554, HEMBB1000573,
                                                           HEMBB1000649.
HEMBB1000652,
              HEMBB1000709, HEMBB10007 49, HEMBB1000790, HEMBB1000827,
              HEMBB1000893, HEMBB1001004, HEMBB1001 008, HEMBB1001047,
HEMBB1000831,
HEMBB1001315,
              HEMBB1001317, HEMBB1001326, HEMBB1001367, HEMBB100 1424,
HEMBB1001436, HEMBB1001458, HEMBB1001535, HEMBB1001565,
                                                           HEMBB1001747.
HEMBB10 01749, HEMBB1001797, HEMBB1001836, HEMBB1001863, HEMBB1001875,
HEMBB1001911, HEMBB1 001922, HEMBB1001925, HEMBB1001944, HEMBB1001983,
              HEMBB1001997, HEMBB 1002092, HEMBB1002247, HEMBB1002266,
HEMBB1001996,
HEMBB1002387, HEMBB1002425, HEMBB1002458, HEMB B1002522, HEMBB1002534,
HEMBB1002582.
              HEMBB1002596, HEMBB1002617, HEMBB1002702, MAMMA1000043,
MAMMA1000092, MAMMA1000129, MAMMA1000198, MAMMA1000221, MAMMA1000307
MAMMA1000331. MAMMA1000360. MAMMA1000402. MAMMA1000414. MAMMA1000500.
MAMMA100052 2, MAMMA1000576, MAMMA1000594, MAMMA1000597, MAMMA1000720,
MAMMA1000775, MAMMA10007 78, MAMMA1000798, MAMMA1000862, MAMMA1000876,
MAMMA1000931, MAMMA1000940, MAMMA1000 941, MAMMA1000975, MAMMA1001038,
MAMMA1001186, MAMMA1001220, MAMMA1001256, MAMMA100 1274, MAMMA1001341,
MAMMA1001397, MAMMA1001420, MAMMA1001547, MAMMA1001670, MAMMA10 01679,
MAMMA1001711, MAMMA1001745, MAMMA1001760, MAMMA1001769, MAMMA1001815, MAMMA1
001907, MAMMA1002056, MAMMA1002078, MAMMA1002093, MAMMA1002125, MAMMA1002132,
MAMMA 1002145, MAMMA1002250, MAMMA1002311, MAMMA1002411, MAMMA1002498,
MAMMA1002571, MAMM A1002701, MAMMA1002727, MAMMA1002728, MAMMA1002746,
MAMMA1002764, MAMMA1002765, MAM MA1002820, MAMMA1002830, MAMMA1002909,
MAMMA1002941, MAMMA1002973, MAMMA1003004, MA MMA1003007, MAMMA1003039,
MAMMA1003089.
NT2RM4000086,
               NT2RM4000265,
                              NT2RM4000414,
                                             NT2RM4000779,
                                                            NT2RM4000855,
                NT2RM4001313, NT2RM4001437,
NT2RM4001160
                                             NT2RM4001754,
                                                           NT2RM4001953,
NT2RM4001984,
              NT2RP200007 7, NT2RP2000183,
                                             NT2RP2000420,
                                                            NT2RP2000678,
NT2RP2000715,
              NT2RP2000842, NT2RP20009 70.
                                             NT2RP2001149,
                                                            NT2RP2001226.
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- NT2RP2001295, NT2RP2001347, NT2RP2001569. NT2RP2001 663. NT2RP2001936 NT2RP2002041, NT2RP2002172, NT2RP2002219, NT2RP2002316, NT2RP200 2546, NT2RP2002591, NT2RP2002643, NT2RP2002741, NT2RP2002750, NT2RP2002778,
- NT2RP20 02857, NT2RP2003000, NT2RP2003073, NT2RP2003237. NT2RP2003394, NT2RP2003517, NT2RP2 003668, NT2RP2003988, NT2RP2004232, NT2RP2004523, NT2RP2004736, NT2RP2004767, NT2RP 2004775, NT2RP2004961, NT2RP2004962, NT2RP2004982, NT2RP2005407, NT2RP2005726, NT2R P2006258. NT2RP2006261,
- NT2RP2006454. NT2RP3000055, NT2RP3000233, NT2RP3000341, NT2 RP3000418, NT2RP3000451, NT2RP3000561, NT2RP3000582, NT2RP3001281, NT2RP3001339, NT 2RP3001340, NT2RP3001383, NT2RP3001432, NT2RP3001580, NT2RP3001589, NT2RP3002004, N T2RP3002173, NT2RP3003133, NT2RP3003346. NT2RP3003403.
- NT2RP3003576, NT2RP3003625, NT2RP3003665, NT2RP3003800, NT2RP3003828, NT2RP3004070, NT2RP3004470, NT2RP4000023 , NT2RP4000035, NT2RP4000102, NT2RP4000167, NT2RP4000214, NT2RP4000218, NT2RP400042 4, NT2RP4000915, NT2RP4002075,
- 20 OVARC1000085, OVARC1000092, OVARC1000145, OVARC1000414, OVARC1000496, OVARC1000526 , OVARC1000948, OVARC1001011, OVARC1001600, OVARC1001805, OVARC1001813, OVARC100184 6,
- PLACE1000540. PLACE1000599, PLACE1001088. PLACE1001377, PLACE1001440, 25 PLACE1001517 PLACE1001672, PLACE1001756, PLACE1002157, PLACE1002205, PLACE1002259, PLACE100239 PLACE1002477, PLACE1002583, PLACE1002968, PLACE1003238, PLACE1003566, PLACE10035 93, PLACE1003618. PLACE1004274. PLACE1004716. PLACE1004815, PLACE1004773, PLACE1004 979, PLACE1005052,
- PLACE1005086, PLACE1005128, PLACE1005176, PLACE1005467, PLACE100 5639, PLACE1005850, PLACE1006003, PLACE1006017, PLACE1006288, PLACE1006371, PLACE10 06629, PLACE1007478, PLACE1008330, PLACE1008584, PLACE1008851, PLACE1008941, PLACE1 009039, PLACE1009493, PLACE1009539, PLACE1009637, PLACE1009947.
- 35 PLACE1010231, PLACE 1010562, PLACE1010579, PLACE1010739. PLACE1010802, PLACE1010896, PLACE1011032, PLAC E1011185, PLACE1011452. PLACE1011465, PLACE1011520. PLACE1011567. PLACE1011719, PLA CE2000011, PLACE2000017, PLACE2000061, PLACE2000187, PLACE2000216, PLACE2000335, PL ACE2000347,
- PLACE2000366, PLACE2000394, PLACE2000398, PLACE2000425, PLACE2000450, PLACE2000477, PLACE3000119, PLACE3000207, PLACE3000230, PLACE3000271, PLACE3000373, PLACE3000399, PLACE3000401, PLACE3000406, PLACE4000247, PLACE4000320, PLACE4000367, PLACE4000401,
- THYRO1000111. THYRO1000187, THYRO1000484, THYRO1000596, THYRO1000625, THYRO1000815 THYRO1000865, THYRO1001003, THYRO1001031, THYRO1001133, THYRO1001401, THYRO100142 6, THYRO1001434, THYRO1001559, THYRO1001570, THYRO1001706, THYRO1001746, THYRO10017 72, THYRO1001907, Y79AA1000033,
- ⁵⁰ Y79AA1000346, Y79AA1000805, Y79AA1001692, Y79AA1002 220.

Group (5): 24 clones

The following 24 clones having low score in the ATGpr1 were judged to be full-length since their ATGpr2 score was 0.3 or higher (Table 11). The clones were novel clones, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

HEMBA1000622, HEMBA1000749, HEMBA1000876, HEMBA1001226, HEMBA1001391, HEMBA1002742 , HEMBA1003908, HEMBA1004000, HEMBB1000336, MAMMA1000356, MAMMA1000883, MAMMA100159 0, NT2RP2000289, NT2RP2001467, PLACE1002853, PLACE1003420, PLACE1004836, PLACE10049 13, PLACE1006795, PLACE1008181, PLACE1008715, PLACE4000344, THYRO1000129, THYRO1001 321,

Group (6): 65 clones

- The following 65 clones having low scores in both the ATGpr1 and ATGpr2 (Table 11) were judged to be full-length, since both scores were still the maximum in a cluster compared with those of the other clones(at least 2 clones or more) in the same cluster. The clones were novel clones, in which at least either of the 5'-end or 3'-end sequence is not identical to any of human ESTs.
 - HEMBA1000604, HEMBA1000673, HEMBA1001024, HEMBA1001026, HEMBA1001734, HEMBA1001784, HEMBA1001808, HEMBA1003902, HEMBA1004164, HEMBA1004909, HEMBA1005232, HEMBA100557, HEMBA1006461, HEMBA1006695, HEMBB1001119,
- HEMBH001337, HEMBH001536, HEMBB10018 68, HEMBB1002045, HEMBB1002579,
 MAMMA1000444, MAMMA1000761, MAMMA1000943, MAMMA1001 820, MAMMA1002360,
 NT2RM4000366, NT2RM4001856, NT2RM4002390, NT2RP2000108, NT2RP2000257.
- NT2RP2001506. NT2RP2002047. NT2RP2002066, NT2RP2002475, NT2RP2004400. 25 NT2RP2004587, NT2RP200528 9, NT2RP2005694, NT2RP3001898, NT2RP3003264, NT2RP3003433. NT2RP3003842, OVARC10012 40, PLACE1001323, PLACE1002227, PLACE1002500, PLACE1002604, PLACE1002772, PLACE1003 478, PLACE1004681. PLACE1005108, PLACE1005932, PLACE1006318, PLACE1006368, PLACE100 6506,
- PLACE1006904, PLACE1007557, PLACE1007877, PLACE1009048, PLACE1011109, PLACE10 11643, PLACE4000548, THYRO1000279, Y79AA1000410, Y79AA1002103,

Group (7): 32 clones

- The following 32 clones having low scores in both the ATGpr1 and ATGpr2 (Table 11) were judged to be full-length, since both scores were still the maximum in a cluster compared with those of the other clones(at least 2 clones or more) in the same cluster. The clones were novel clones, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.
 - HEMBA1000251, HEMBA1001803, HEMBA1001918, HEMBA1002257, HEMBA1003064, HEMBA1003714 , HEMBA1004405, HEMBA1005508, HEMBB1000054, HEMBB1001142, MAMMA1000175, MAMMA100116 2, MAMMA1002972, NT2RM4000425, NT2RP2004512,
- 45 NT2RP2005531, NT2RP2005942, NT2RP20065 54, NT2RP3001007, NT2RP3001318, OVARC1000017, OVARC1000068, OVARC1000486, PLACE1001 705, PLACE1002319, PLACE1007743, PLACE1007829, PLACE1008630, PLACE1009925, PLACE101 1492, PLACE1011749, THYRO1000793,
- The following 117 clones, selected by assembling the sequence of the other clones in the same cluster and human ESTs, have high fullness ratio. The clones were classified into the following group (8) and (9).
- HEMBA1001323, HEMBA1001330, HEMBA1001712, HEMBA1001820, HEMBA1002204, HEMBA1002349, HEMBA1002538, HEMBA1003309, HEMBA1003939, HEMBA1004015, HEMBA1004295, HEMBA100467 2, HEMBA1004865, HEMBA1005251, HEMBA1006158, HEMBA1006676, HEMBA1006779, HEMBA10072 88, HEMBB1000218, HEMBB1000272,

- HEMBB1000399, HEMBB1000491, HEMBB1000996, HEMBB1001 114, HEMBB1001850, HEMBB1002015, MAMMA1000287, MAMMA1001683, MAMMA1001686, MAMMA100 2612,
- NT2RM2000609, NT2RM4002438, NT2RM4002567, NT2RP2000270, NT2RP2000758, 5 NT2RP2001290, NT2RP2001526, NT2RP2002124, NT2RP2002736, NT2RP2002753, NT2RP2003456, NT2RP200372 7, NT2RP2003871, NT2RP2003968, NT2RP2004321, NT2RP2004580, NT2RP20052 93, NT2RP2004412, NT2RP2005476, NT2RP2005753, NT2RP2005815, NT2RP2005841, NT2RP2005857, NT2RP2006 393, NT2RP2006467,
- NT2RP3000109, NT2RP3000449, NT2RP3001245, NT2RP3001634, NT2RP300 2056, NT2RP3002810, NT2RP3002955, NT2RP3003032, NT2RP3003138, NT2RP3003500, NT2RP30 03819, NT2RP4000078, NT2RP4000515, NT2RP4000517, NT2RP4001407, NT2RP4001889, NT2RP4 002905,
- OVARC1000071, OVARC1001883, PLACE1000292, PLACE1001007, PLACE1001395, PLACE1001691 , PLACE1001746, PLACE1001748, PLACE1001845, PLACE1002066, PLACE1003373, PLACE100390 0, PLACE1004118, PLACE1004256, PLACE1004284, PLACE1004336, PLACE1004506, PLACE10049 34, PLACE1005077, PLACE10051111,
- PLACE1005409, PLACE1005730, PLACE1006076, PLACE1006 360, PLACE1006470, PLACE1006760, PLACE1006867, PLACE1007045, PLACE1007111, PLACE100 7807, PLACE1008080, PLACE1008244, PLACE1008369, PLACE1008405, PLACE1008426, PLACE10 08621, PLACE1009020, PLACE1009621, PLACE1010089, PLACE1010270, PLACE3000276,
- ²⁵ THYRO1 000805, THYRO1001365, THYRO1001673, Y79AA1001848,

Group (8): 36 clones

- Among the 117 clones described above, the following 36 clones were novel clones, in which at least either of the 5'-end or 3'-end sequence, or both of them were not identical to any of human ESTs.
 - HEMBA1001330, HEMBA1001712, HEMBA1001820, HEMBA1002204, HEMBA1002349, HEMBA1003939 , HEMBA1004295, HEMBA1004865, HEMBA1006779, HEMBB1000218,
- 35 HEMBB1000491, HEMBB100099 6, HEMBB10011114, MAMMA1000287, MAMMA1001686, MAMMA1002612, NT2RP2002736, NT2RP20037 27, NT2RP2004580, NT2RP2005841, NT2RP2006393, PLACE1002066, PLACE1003373, PLACE1003 900, PLACE1004118, PLACE1004336, PLACE1004506, PLACE1004934, PLACE1005409, PLACE100 5730,
- 40 PLACE1006470, PLACE1008080, PLACE3000276, THYRO1000805, THYRO1001365, THYRO10 01673,

Group (9): 81 clones

- Among the 117 clones described above, the following 81 clones were the clones in which the number of the identical human ESTs for the 5'-end sequence is not more than 20 (except clones in which at least either of the 5'-end or 3'-end sequence, or both of them are not identical to any of human ESTs, and clones in which the number of the identical human ESTs for both the 5'-
- end and 3'-end sequences is 1 or higher and not more than 5).
 HEMBA1001323, HEMBA1002538, HEMBA1003309, HEMBA1004015, HEMBA1004672,
 HEMBA1005251 , HEMBA1006158, HEMBA1006676, HEMBA1007288, HEMBB1000272,
 HEMBB1000399, HEMBB100185 0, HEMBB1002015, MAMMA1001683, NT2RM2000609,
- NT2RM4002438, NT2RM4002567, NT2RP20002 70, NT2RP2000758, NT2RP2001290, NT2RP2001526, NT2RP2002124, NT2RP2002753, NT2RP2003 456, NT2RP2003871, NT2RP2003968, NT2RP2004321, NT2RP2004412, NT2RP2005293, NT2RP200 5476,

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Group (10): 938 clones

The following 938 clones having low scores in both the ATGpr1 and ATGpr2 were judged to be full-length according to the fullness ratio shown in Table 4. The clones were novel clones, in which at least the 5'-end sequence was not identical to any of human ESTs.

HEMBA100042, HEMBA1000213, HEMBA1000264, HEMBA1000355, HEMBA1000366, HEMBA1000411, HEMBA1000422, HEMBA1000428, HEMBA1000434, HEMBA1000442, HEMBA1000464, HEMBA100054 0, HEMBA1000563, HEMBA1000569, HEMBA1000655, HEMBA1000662, HEMBA1000702, HEMBA10007 05, HEMBA1000722, HEMBA1000747, HEMBA1000769, HEMBA1000773, HEMBA1000774, HEMBA1000 827, HEMBA1000843, HEMBA1000852, HEMBA1000870, HEMBA1000908, HEMBA1000985, HEMBA100 0991, HEMBA1001017, HEMBA1001019, HEMBA1001094, HEMBA1001121, HEMBA1001172, HEMBA10 01265, HEMBA1001327, HEMBA1001375, HEMBA1001432, HEMBA1001454,

- 30 HEMBA10 01265, HEMBA1001327, HEMBA1001375, HEMBA1001432, HEMBA1001454, HEMBA1001463, HEMBA1 001515, HEMBA1001517, HEMBA1001557, HEMBA1001566, HEMBA1001585, HEMBA1001589, HEMBA 1001608, HEMBA1001675, HEMBA1001678, HEMBA1001681, HEMBA1001718, HEMBA1001761, HEMB A1001945, HEMBA1001960,
- 35 HEMBA1001964, HEMBA1001991,
 - HEMBA1002003, HEMBA1002008, HEMBA1002039, HEMBA1002100, HEMBA1002113, HEMBA1002139 , HEMBA1002144, HEMBA1002153, HEMBA1002226, HEMBA1002253, HEMBA1002270, HEMBA100232 1, HEMBA1002328, HEMBA1002337, HEMBA1002381,
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- ³⁰ Y79AA1000538, Y79AA1000802, Y79AA1000969, Y79AA1001061, Y79AA1 001167, Y79AA1001384, Y79AA1001594, Y79AA1001875, Y79AA1002211,

Group (11): 228 clones

- The following 228 clones having low scores in both the ATGpr1 and ATGpr2 were judged to be full-length according to the fullness ratio shown in Table 7. The clones were a novel clone, in which at least the 3'-end sequence was not identical to any of human ESTs.
- HEMBA1000111, HEMBA1000243, HEMBA1000338, HEMBA1000390, HEMBA1000418, HEMBA1000459 , HEMBA1000518, HEMBA1000791, HEMBA1000974, HEMBA1001007, HEMBA100109 9, HEMBA1001123, HEMBA1001294, HEMBA1001299, HEMBA1001442, HEMBA1001861, HEMBA10019 42, HEMBA1001979, HEMBA1002160, HEMBA1002162, HEMBA1002166, HEMBA1002389, HEMBA1002 498, HEMBA1002592,
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40 THYRO1001595, THYRO 1001745.

EXAMPLE 12

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Homology search using the 5'-end and 3'-end sequences of the selected clones.

The 5' -end sequences of the selected 5547 clones were used for the homology search of the SwissProt, and both the 5' -end 3' -end sequences were used for the search of the GenBank and UniGene (ref. the result of the search of the SwissProt, GenBank (except ESTs and STSs), and UniGene (Human) was attached).

11964, PLACE2000136, PLACE2000264, PLACE2000305, PLACE2000379, PLACE3000220, PLACE4 000250, THYRO1000092,THYRO1000471, THYRO1000852, THYRO1001534,

Each search result is shown in the last part of this SPECIFICATION by arranging each item in the following format.

	5'-end sequence	3'-end sequence
Swiss-Prot	Homology search result 1	*****
GenBank	Homology search result 2	Homology search result 3
UniGene	Homology search result 4	Homology search result 5
	GenBank	Swiss-Prot Homology search result 1 GenBank Homology search result 2

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According to the top hit data, at least 1430 clones were predicted to encode a protein belonging to any of the categories, secretory or membrane protein, glycoprotein, protein associated with signal transduction, protein associated with transcription, protein associated with diseases, enzyme or protein associated with metabolism, protein associated with cell division or cell proliferation, protein associated with cytoskeleton, protein associated with RNA synthesis, nuclear protein, protein associated with protein synthesis or transport, protein associated with cellular defense, or protein associated with development or growth. Among the clones predicted belonging to any of the categories, 1001 clones were estimated to have a relatively high homology with the known proteins or genes in the same category. In addition, 429 clones were estimated to have a relatively low homology with the known proteins in the same category.

Herein, the term "relatively high homology" is defined as having 60% or more identity and the P-value 10⁻¹⁰à or less in comparison with known sequences in the SwissProt database, or 64% or more identity and the P-value 10⁻¹⁵à or less in comparison with those in the GenBank and UniGene databases (see the attached list). Also, the term "relatively low homology" is defined as not fulfilling the requirements to be "relatively high homology", but still having the scores, 25% or more identity and the P-value 10⁻⁶à or less, using the sequence having 55 nucleotides or more, in comparison with known sequences in the SwissProt database (see the attached list). The P-value is a score obtained statistically by taking into account the probability of occurrence of the similarity between two sequences. In general, the smaller P-value reflects the higher similarity (Altschul S.F., Gish W., Miller W., Myers E.W., and Lipman D.J. (1990) "Basic local alignment search tool" J.Mol. Biol., 215: 403-410; Gish W., and States D.J. (1993) "Identification of protein coding regions by database similarity search" Nature Genet. 3: 266-272).

The clones predicted to encode a protein in the category of secretory protein or membrane protein have the keywords, "signal", "transmembrane", "membrane", "extracellular matrix", "receptor", "G-protein coupled receptor", "ionic channel", "voltage-gated channel", "calcium channel", "cell adhesion", "collagen", or "connective tissue", or descriptions from which the clone can be predicted to be a secretory or membrane protein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

The clones predicted to encode a protein in the category of glycoprotein have the keywords, "glycoprotein", or descriptions from which the clone can be predicted to be a glycoprotein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

The clones predicted to encode a protein in the category of proteins associated with signal transduction have the keywords, "serine/threonine-protein kinase", "tyrosine-protein kinase", "SH3 domain", or "WD repeat", or descriptions from which the clone can be predicted to be a protein associated with signal transduction (such as "ADP-ribosylation factor"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or

UniGene using the 5'-end sequence and 3'-end sequence.

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The clones predicted to encode a protein in the category of proteins associated with transcription have the keywords, "transcription regulation", "zinc finger", or "homeobox", or descriptions from which the clone can be predicted to be a protein associated with transcription, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

The clones predicted to encode a protein in the category of proteins associated with diseases are the clones in which the top hit data of the SwissProt using the 5'-end sequence, or the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence is a gene or protein that is deposited in the Online Mendelian Inheritance in Man (OMIM) database, which is a database of human genes and diseases, or the top hit data has descriptions from which the clone can be predicted to be a protein associated with diseases.

The clones predicted to encode a protein in the category of enzyme or proteins associated with metabolism are the clones in which the top hit data of the SwissProt using the 5'-end sequence, or the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence is a gene or protein with E.C.No. (Enzyme commission number), or the top hit data has descriptions from which the clone can be predicted to be an enzyme or protein associated with metabolism (such as "metabolism", "oxidoreductase", or "lipid").

The clones predicted to encode a protein in the category of proteins associated with cell division or cell proliferation have the keywords, "cell division", "cell cycle", "mitosis", or "chromosomal protein", or descriptions from which the clone can be predicted to be a protein associated with cell division or cell proliferation (such as "histone", "cell growth", or "apoptosis"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

The clones predicted to encode a protein in the category of proteins associated with cytoskeleton have the keywords, "structural protein", "cytoskeleton", "actin-binding", or "microtubules", or descriptions from which the clone can be predicted to be a protein associated with cytoskeleton, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

The clones predicted to encode a protein in the category of proteins associated with RNA synthesis include the above clones predicted to be a protein associated with transcription, and also the clones which have the keywords, "RNA splicing", or "RNA processing", or descriptions from which the clone can be predicted to be a protein associated with RNA synthesis (such as "RNA helicase", "polyadenylation", or "RNA transport"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

The clones predicted to encode a protein in the category of nuclear protein include the above clones predicted to be a protein associated with transcription, and also the clones which have the keyword, "nuclear protein", or descriptions from which the clone can be predicted to be a nuclear protein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

The clones predicted to encode a protein in the category of proteins associated with protein synthesis or transport have the keywords, "translation regulation", "protein biosynthesis", "amino-acid biosynthesis", "ribosomal protein", or "protein transport", or descriptions from which the clone can be predicted to be a protein associated with protein synthesis or transport

(such as "signal recognition particle", "ubiquitin", "proteosome", or "protease"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

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The clones predicted to encode a protein in the category of proteins associated with cellular defense have the keywords, "heat shock", "chaperone", "DNA repair", or "DNA damage", or descriptions from which the clone can be predicted to be a protein associated with cellular defense, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

The clones predicted to encode a protein in the category of proteins associated with development or growth have the keyword, "developmental protein", or descriptions from which the clone can be predicted to be a protein associated with development or growth, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

The following 1430 clones were predicted to encode a protein belonging to any of the categories, secretory or membrane protein, glycoprotein, protein associated with signal transduction, protein associated with transcription, protein associated with diseases, enzyme or protein associated with metabolism, protein associated with cell division or cell proliferation, protein associated with cytoskeleton, protein associated with RNA synthesis, nuclear protein, protein associated with protein synthesis or transport, protein associated with cellular defense, or protein associated with development or growth.

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       Among the clones, 1001 clones were estimated to have a relatively high homology with the
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THYRO1000387.
                 THYRO1000471.
                                 THYRO1000501.
                                                  THYRO1000585,
                                                                  THYRO1000666.
THYRO1000783,
                 THYRO1000926,
                                 THYRO1000974,
                                                  THYRO1001100,
                                                                  THYRO1001121,
THYRO1001189.
                 THYRO1001406,
                                 THYRO1001458.
                                                  THYRO1001595,
                                                                  THYRO1001605,
THYRO1001617,
                 THYRO1001671,
                                  Y79AA1000037,
                                                  Y79AA1000059,
                                                                   Y79AA1000214,
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	Y79AA1000346,	Y79AA1000349,	Y79AA1000469,	Y79AA1000560	Y79AA1000627,
	Y79AA1000705,	Y79AA1000734,	Y79AA1000752,	Y79AA1000784	, Y79AA1000833,
	Y79AA1000966,	Y79AA1000968,	Y79AA1000985,	Y79AA1001105	Y79AA1001236,
5	Y79AA1001299,	Y79AA1001391,	Y79AA1001533,	Y79AA1001548	, Y79AA1001679,
	Y79AA1001827,	Y79AA1001866,	Y79AA1002093,	Y79AA1002258	, Y79AA1002361,
	Y79AA1002416,				
	The following	429 clones were es	stimated to have a	relatively low homo	logy with the known
10	proteins in the sa	me category.			
	- HEMBA1000012.	HEMBA1000185.	HEMBA1000488.	HEMBA1000491,	HEMBA1000569.
	HEMBA1000752,	inamonito do 100,	maiom voo roo,	industrious 131,	inamonii oo o
	manditivoo to 2,				
15	HEMBA1001137,	HEMBA1001407.	HEMBA1001476.	HEMBA1001744.	HEMBA1001800.
	· · · · · · · · · · · · · · · ·	HEMBA1001809, HE			
	,				
	HEMBA1002035.	HEMBA1002102,	HEMBA1002150,	HEMBA1002212.	HEMBA1002495,
20		HEMBA1002609, HE		•	
	,	,	,	,	
25					
23					

		EP 1 074 617 A2
	HEMBA1003071, HEMBA1003291,	HEMBA1003077, HEMBA1003136, HEMBA1003179, HEMBA1003235, HEMBA1003418,
5	HEMBA1003568, HEMBA1003953,	НЕМВА1003598, НЕМВА1003680, НЕМВА1003684, НЕМВА1003836,
10	HEMBA1004276,	HEMBA1004408, HEMBA1004669, HEMBA1004734,
	HEMBA1005475,	НЕМВА1005595, НЕМВА1005737, НЕМВА1005815, НЕМВА1005931,
	HEMBA1006347,	HEMBA1006474, HEMBA1006521, HEMBA1006562, HEMBA1006652,
15	HEMBA1006737,	HEMBA1006914,
	HEMBA1007045,	НЕМВА1007121, НЕМВА1007300, НЕМВА1007301,
20	HEMBB1000119,	HEMBB1000217, HEMBB1000317, HEMBB1000632, HEMBB1000789,
	HEMBB1001011,	HEMBB1001177, HEMBB1001234, HEMBB1001282, HEMBB1001288,
25	HEMBB1002050,	HEMBB1002092,
	MAMMA1000085,	MAMMA1000672, MAMMA1000841, MAMMA1000897, MAMMA1001008,
	MAMMA1001030,	MAMMA1001041,
30	MAMMA1002143,	MAMMA1002573, MAMMA1002619, MAMMA1002636, MAMMA1002650,
	•	MAMMA1002881, MAMMA1002937,
35	NT2RM1000118,	NT2RM1000153, NT2RM1000186, NT2RM1000187, NT2RM1000377,
	NT2RM1000648,	•
	NT2RM1000850,	NT2RM1000882, NT2RM1000898,
40	NT2RM1001059,	NT2RM1001072, NT2RM1001139,
	NT2RM2000363,	NT2RM2000371, NT2RM2000402, NT2RM2000577, NT2RM2000599,
45	NT2RM2000623,	NT2RM2000714,
	NT2RM2001131,	NT2RM2001221, NT2RM2001324, NT2RM2001345,
	NT2RM2001588,	NT2RM2001635, NT2RM2001670, NT2RM2001700, NT2RM2001730,
50	NT2RM2001785,	NT2RM2001797, NT2RM2001823, NT2RM2001936, NT2RM2001989,
	NT2RM2002580,	
	NT2RM4000030,	NT2RM4000085, NT2RM4000155, NT2RM4000156, NT2RM4000169,
55	NT2RM4000191,	NT2RM4000215, NT2RM4000354, NT2RM4000421, NT2RM4000496,
	NT2RM4000734,	NT2RM4000751,

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	NT2RM4001032, NT2RM4001444,	NT2RM4001140,	NT2RM4001316,	NT2RM4001340,	NT2RM4001410,
5	NT2RM4001611, NT2RM4001930,	NT2RM4001629, NT2RM4001979, NT2	NT2RM4001746, RM4001987,	NT2RM4001776,	NT2RM4001783,
10	NT2RM4002062,	NT2RM4002145, NT2	RM4002189, NT2RN	14002226, NT2RM40	02409,
	NT2RM4002527,	NT2RM4002571, NT2	RM4002623,		
15	NT2RP1000111, NT2RP1000460,	NT2RP1000130, NT2RP1000470, NT2	NT2RP1000333, RP1000493,	NT2RP1000348,	NT2RP1000363,
20	NT2RP1000547, NT2RP1001543,	NT2RP1000609,	NT2RP1001033,	NT2RP1001080,	NT2RP1001313,
	NT2RP2000006, NT2RP2000448,	NT2RP2000133,	NT2RP2000161,	NT2RP2000183,	NT2RP2000257,
25	NT2RP2000710, NT2RP2001174,	NT2RP2000764, NT2RP2001839, NT2	NT2RP2000812, RP2001900,	NT2RP2000932,	NT2RP2001127,
30	NT2RP2002025, NT2RP2002618,	NT2RP2002058, NT2RP2002710, NT2	NT2RP2002185, RP2002980,	NT2RP2002442,	NT2RP2002595,
35	NT2RP2003125, NT2RP2003564,	NT2RP2003277, NT2RP2003706,	NT2RP2003286,	NT2RP2003293,	NT2RP2003506,
	NT2RP2004098, NT2RP2005037,	NT2RP2004187, NT2RP2005038,	NT2RP2004568, NT2RP2005204,	NT2RP2004791, NT2RP2005344,	NT2RP2005022, NT2RP2005465.
40	NT2RP2005496, NT2RP2006275,	NT2RP2005531,	NT2RP2005690,	NT2RP2005763,	NT2RP2005767,
45	NT2RP3000046, NT2RP3000527, NT2RP3000994,	NT2RP3000047, NT2RP3000590,	NT2RP3000361, NT2RP3000603,	NT2RP3000397, NT2RP3000759,	NT2RP3000418, NT2RP3000868,
50	NT2RP3001216, NT2RP3001426,	NT2RP3001274, NT2RP3001427,	NT2RP3001355,	NT2RP3001399,	NT2RP3001407,
-	NT2RP3001587, NT2RP3002399,	NT2RP3001708, NT2RP3002663, NT2	NT2RP3002056, RP3002785, NT2RP	NT2RP3002273, 23002909,	NT2RP3002377,
55	NT2RP3003061, NT2RP3003825,	NT2RP3003500, NT2RP3003831,	NT2RP3003672,	NT2RP3003701,	NT2RP3003716,

•	NT2RP3004209, NT2RP3004507,	NT2RP3004253, NT2RP3004454, NT2RP3004472, NT2RP30 NT2RP3004569, NT2RP3004578, NT2RP3004617, NT2RP3004669,	04475,
5	•		
	NT2RP4000210,	NT2RP4000212, NT2RP4000259, NT2RP4000312, NT2RP40	00370,
	NT2RP4000417,	NT2RP4000449, NT2RP4000457, NT2RP4000498, NT2RP40	00528,
	NT2RP4000879,	NT2RP4000925,	
10			
	NT2RP4001041,	NT2RP4001095, NT2RP4001122, NT2RP4001126, NT2RP40	
	NT2RP4001149,	NT2RP4001150, NT2RP4001174, NT2RP4001207, NT2RP40	01219,
46	NT2RP4001372,	NT2RP4001375,	
15	NT2RP4001551,	NT2RP4001592, NT2RP4001638, NT2RP4001725, NT2RP40	01700
	NT2RP4001823,		
	NT2RP4001946,	NT2RP4001838, NT2RP4001849, NT2RP4001893, NT2RP40 NT2RP4002408,	01927,
20		11324 2002 1009	
20	NT2RP5003477,	NT2RP5003522, NT2RP5003524,	_
		•	
	OVARC1000004,	OVARC1000087, OVARC1000139, OVARC1000309, OVARC10	00543,
25	OVARC1000746,	OVARC1000846, OVARC1000876, OVARC1000885, OVARC1000912,	
	OVARC1001180,	OVARC1001329, OVARC1001476, OVARC1001555, OVARC10	01610,
	OVARC1001762,	OVARC1001880, OVARC1001942, OVARC1002050, OVARC1002138,	
30	PLACE1000066,	PLACE1000142, PLACE1000420, PLACE1000547, PLACE10	00500
	PLACE1000863,	PLACE1000142, PLACE1000420, PLACE1000547, PLACE10	00583,
	121002000000,		j
	PLACE1001387,	PLACE1001632, PLACE1001692, PLACE1001989,	
35	ŕ	, , ,	
	PLACE1002450,	PLACE1002571, PLACE1002722, PLACE1002775, PLACE10	02811,
	PLACE1002991,		,
40		•	
40	PLACE1003190,	PLACE1003258, PLACE1003420, PLACE1003493, PLACE10	03605,
	PLACE1003669,	PLACE1003704, PLACE1003915,	
	DI ACRICO 4102	DI ACCIONATOR DI AGRICOLATE	
45	PLACE1004197,	PLACE1004428, PLACE1004471, PLACE1004629, PLACE10	04751,
	PLACE1004902,		
	PI 4/F1005287	PLACE1005557, PLACE1005611, PLACE1005763, PLACE1005890,	
	I Linchitotopor,	TENCETOUSSET, FENCETOUSSET, FENCETOUSSES, FENCETOUSSES,	
50	PLACE1006157.	PLACE1006469, PLACE1006731,	•
	,		
	PLACE1007238,	PLACE1007375, PLACE1007488, PLACE1007544, PLACE100	07598.
	•	PLACE1008244, PLACE1008275, PLACE1008947,	,
55		•	
	PLACE1009094,	PLACE1009319, PLACE1009622, PLACE1009721, PLACE100	09845,
_			-

	PLACE1009861,				
5	PLACE1010083, PLACE1010310, PLACE1011263,	PLACE1010089, PLACE1010833, PLACE1011371, PLA	PLACE1010106, PLACE1010960, ACE1011492,	PLACE1010152, PLACE1011056,	PLACE1010194, PLACE1011219,
10	PLACE2000034, PLACE2000458,	PLACE2000164,	PLACE2000274,	PLACE2000399,	PLACE2000411,
15	PLACE3000148, PLACE3000373,	PLACE3000156,	PLACE3000242,	PLACE3000322,	PLACE3000353,
	PLACE4000009, PLACE4000558,	PLACE4000014, PLACE4000581, PL	PLACE4000259, ACE4000590, PLAC	PLACE4000431, E4000612,	PLACE4000522,
20	SKNMC1000011,				
0.5	THYR01000072,	THYR01000852, TH	YR01000934, THYR	01000951,	·
25	Y79AA1000258,	•	•		Y79AA1000962;
	Y79AA1001048, Y79AA1001692,	Y79AA1001233, Y79AA1001848, Y79	Y79AA1001394, 9AA1001874, Y79A	=	Y79AA1001613,
30					
	Y79AA1002027, Y79AA1002482,	Y79AA1002139,	Y79AA1002208,	Y79AA1002209,	Y79AA1002472,

(1) Among the 1430 clones, the following 259 clones were predicted to encode a protein in the category of secretory protein or membrane protein (including the clones belonging to plural categories).

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HEMBA1001407 , HEMBA1001476, HEMBA1001569, HEMBA1001678, HEMBA1001804, HEMBA1002150, HEMBA100254 7, HEMBA1002555, HEMBA1002688, HEMBA1002716, HEMBA1002999, HEMBA1003071, HEMBA10030 77, HEMBA1003538, HEMBA1003598, HEMBA1003690, HEMBA1004207, HEMBA1004356, HEMBA1004 389, HEMBA1004756, HEMBA1005338, HEMBA1005367, HEMBA1005576, HEMBA1005699, HEMBA100 5963, HEMBA1005990, HEMBA1006310, HEMBA1006344, HEMBA1006562, HEMBA1006976, HEMBA100530, HEMBA1007301, HEMBB1000037, HEMBB1000119, HEMBB1000317, HEMBB1000530, HEMBB1 000915, HEMBB1001112, HEMBB1001119, HEMBB1001151, HEMBB1001282, HEMBB1001331, HEMBB 1002042, HEMBB1002092, HEMBB1002193, HEMBB1002103, MANAMALO00424, MANAMALO00673, M

HEMBA1000005, HEMBA1001017, HEMBA1001059, HEMBA1001071, HEMBA1001351,

HEMBB1002600, MAMMA1000103, MAMMA1000424, MAMM A1000672, MAMMA1000734, MAMMA1000897, MAMMA1001008, MAMMA1001030, MAMMA1001041, MAM MA1001075, MAMMA1001080, MAMMA1001751, MAMMA1001754, MAMMA1002143, MAMMA1002219, MA

- MMA1002485, MAMMA1002636, MAMMA1002938, MAMMA1003104, NT2RM1000199, NT2RM1000355, N T2RM1000430, NT2RM1000648, NT2RM1000742, NT2RM1000770, NT2RM1000811, NT2RM1000833, NT2RM1000867, NT2RM1000882, NT2RM1000883, NT2RM1001139, NT2RM2000374, NT2RM2000402 , NT2RM2000422, NT2RM2000566, 5 NT2RM2000623, NT2RR2001324, NT2RM2001499, NT2RM200158 8, NT2RM2001592, NT2RM2001613, NT2RM2001635, NT2RM2001648, NT2RM2001760, NT2RM20021 09, NT2RM4000156. NT2RM2002145, NT2RM4000169. NT2RM4000386, NT2RM4000433, NT2RM4000795, NT2RM4001032, NT2RM4001054, NT2RM4001410, 10 NT2RM4001605, NT2RM4001930, NT2RM4001987, NT2RM4002073, NT2RM4002145, NT2RM4002189, NT2RM4002558, NT2RM400256 5, NT2RM4002571, NT2RM4002594, NT2RP1000130, NT2RP1000326, NT2RP1000363, NT2RP10004 13, NT2RP1000547, 15 NT2RP1000782, NT2RP1000856, NT2RP1001247, NT2RP1001313, NT2RP1001 546, NT2RP1001569, NT2RP2000056, NT2RP2000257, NT2RP2000842, NT2RP2001081, NT2RP200 1991, NT2RP2002025, NT2RP2002193, NT2RP2002256, NT2RP2002312, NT2RP2002479, NT2RP20 02504, NT2RP2003433, NT2RP2003706, NT2RP2003760, 20 NT2RP2004194, NT2RP2004655, NT2RP2 005020, NT2RP2005344, NT2RP2005465, NT2RP2005752, NT2RP2005753, NT2RP2005835, NT2RP 2005933, NT2RP2006275, NT2RP2006565, NT2RP2006571, NT2RP3000324, NT2RP3000739, NT2R P3000919, NT2RP3001140, NT2RP3001355, NT2RP3001676, NT2RP3001708, NT2RP3001727, NT2 25 RP3002659, NT2RP3003059, NT2RP3003101, NT2RP3003203, NT2RP3003212. NT2RP3003242, NT 2RP3003672, NT2RP3003701, NT2RP3003716, NT2RP3003918, NT2RP3004207, NT2RP3004253, N T2RP4000008, NT2RP4000212, NT2RP4000243, NT2RP4000417, NT2RP4000878, NT2RP4000907, NT2RP4000925, NT2RP4000928, 30 NT2RP4001079, NT2RP4001117, NT2RP4001150, NT2RP4001174 , NT2RP4001313, NT2RP4001315, NT2RP4001345, NT2RP4001372, NT2RP4001574, NT2RP400182 3, NT2RP4001966, NT2RP5003524, OVARC100060, OVARC1000326, OVARC1000543, OVARC10006 82, OVARC1000722, OVARC1000912, OVARC1001055, OVARC1001154, 35 OVARC1001261, OVARC1001329, OVARC1001419, OVARC1001506, OVARC1001542, OVARC1001610, OVARC1001713 , OVARC1001802, OVARC1001943, PLACE1000611, PLACE1001092, PLACE1001136, PLACE100129 4, PLACE1002437, PLACE1002500, PLACE1002722, 40 PLACE1002782, PLACE1002816, PLACE10034 20, PLACE1003493, PLACE1003596, PLACE1004104, PLACE1004197, PLACE1004203, PLACE1004 277, PLACE1004629, PLACE1004751, PLACE1005494, PLACE1005804, PLACE1005890, PLACE100 6157, PLACE1007409, PLACE1007852, PLACE1008044, PLACE1008273, PLACE1008309, 45 PLACE10 08603, PLACE1008643, PLACE1009659, PLACE1009670, PLACE1009845, PLACE1009861, PLACE1 011371, PLACE1011492, PLACE1011896, PLACE2000034, PLACE2000341, PLACE2000399, PLACE 2000458, PLACE3000020, PLACE3000322, PLACE3000353, PLACE4000052, PLACE4000269, PLAC E4000522, PLACE4000581, 50 THYRO1000783, Y79AA1000258, Y79AA1000346, Y79AA1001874,
 - (2) The following 111 clones were predicted to encode a protein in the category of glycoprotein (including the clones belonging to plural categories).
- HEMBA1000752, HEMBA1001017, HEMBA1001059, HEMBA1001071, HEMBA1001302, HEMBA1002150, HEMBA1002547, HEMBA1002555, HEMBA1003077, HEMBA1003538, HEMBA1003598, HEMBA100386 6, HEMBA1004356, HEMBA1005338, HEMBA1005576,

HEMBA1005699, HEMBA1006562, HEMBA10069 76, HEMBA1007301, HEMBB1000317, HEMBB1000530, HEMBB1001119, HEMBB1002092, HEMBB1002 193, MAMMA1000672, MAMMA1000897, MAMMA1001030, MAMMA1001388, MAMMA1001771, MAMMA100 2485, 5 MAMMA1002636, MAMMA1002938, NT2RM1000648, NT2RM2000374, NT2RM2000407, NT2RM20 00422, NT2RM2000623, NT2RM2001499, NT2RM2001588, NT2RM2001839, NT2RM2001930, NT2RM4 000156, NT2RM4000233, NT2RM4001410, NT2RM4001987, NT2RM4002145, NT2RM4002189, NT2RM 4002194, NT2RM4002558, NT2RM4002571, NT2RP1000856, NT2RP1001247, NT2RP2000056, NT2R P2001991, NT2RP2002025, 10 NT2RP2002325, NT2RP2002385, NT2RP2003706, NT2RP2005933, NT2 RP3001140, NT2RP3003242, NT2RP3003672, NT2RP3003701, NT2RP3003716, NT2RP3003914, NT 2RP3004253, NT2RP4000212, NT2RP4000417, NT2RP4000925, NT2RP4001149, 15 NT2RP4001150, N T2RP4001219, NT2RP4001345, NT2RP4001372, NT2RP5003522, NT2RP5003524, OVARC1000543, OVARC1000682, OVARC1000722, OVARC1001055, OVARC1001154, OVARC1001506, OVARC1001713 PLACE1002437, PLACE1002722, PLACE1003258, PLACE1003493, PLACE1004197, PLACE100475 1, PLACE1005698, 20 PLACE1005804, PLACE1006157, PLACE1006754, PLACE1008643, PLACE10090 94, PLACE1009861, PLACE1011371, PLACE2000034, PLACE2000062, PLACE2000399, PLACE2000458, PLACE3000009, PLACE3000020, PLACE3000353, PLACE3000373, PLACE4000052 , PLACE4000230, PLACE4000522, PLACE4000581, THYRO1000852, 25 Y79AA1001874.

(3) The following 147 clones were predicted to encode a protein in the category of proteins associated with signal transduction (including the clones belonging to plural categories).

30 HEMBA1000030, HEMBA1000185, HEMBA1000303, HEMBA1000459, HEMBA1000491, HEMBA1000657, HEMBA1000991, HEMBA1001174, HEMBA1001387, HEMBA1001822, HEMBA1001921, HEMBA100221 2, HEMBA1002341, HEMBA1002417, HEMBA1002810, HEMBA1003291, HEMBA1003291, HEMBA100333 14, HEMBA1003560, HEMBA1003615,

HEMBA1003621, HEMBA1003805, HEMBA1004286, HEMBA1004 604, HEMBA1006173,
 HEMBA1006540, HEMBA1006973, HEMBA1007174, HEMBB1000632, HEMBB100 0781,
 HEMBB1001051, HEMBB1001177, HEMBB1001234, HEMBB1001294, HEMBB1001384,
 HEMBB10 02193, HEMBB1002477, HEMBB1002635, MAMMA1001305, MAMMA1002699.

MAMMA1002842, MAMMA1 003057, MAMMA1003113, NT2RM1000118, NT2RM1000186, NT2RM1000772, NT2RM1000850, NT2RM 1001072, NT2RM2000124, NT2RM2000612, NT2RM2001065, NT2RM2001221, NT2RM2001345, NT2R M2001652, NT2RM2001983, NT2RM2002109, NT2RM2002128, NT2RM4000327, NT2RM4000354, NT2 RM4000798,

NT2RM4001203, NT2RM4001411, NT2RM4001412, NT2RM4001582, NT2RM4001629, NT 2RM4001758, NT2RM4002226, NT2RP1000018, NT2RP1000112, NT2RP1000701, NT2RP1000825, N T2RP1001395, NT2RP1001410, NT2RP1001457, NT2RP1001482, NT2RP2002058, NT2RP2002710, NT2RP2003702, NT2RP2003912, NT2RP2004232,

NT2RP2004768, NT2RP2005022, NT2RP2005620 , NT2RP2005890, NT2RP3000299, NT2RP3000403, NT2RP3000742, NT2RP3000845, NT2RP300176 4, NT2RP3002004, NT2RP3002785, NT2RP3002909, NT2RP3003230, NT2RP3003411, NT2RP30034 64, NT2RP3003800, NT2RP3004475, NT2RP3004534, NT2RP4000147, NT2RP4000376,

NT2RP4001122, NT2RP4001375, NT2RP4001644, NT2RP4001725, NT2RP4001760,
 NT2RP4001849, NT2RP4001927, NT2RP4002408, NT2RP5003477, OVARC1000437,
 OVARC1000465, OVARC100064 9, OVARC1000945, OVARC1001476, OVARC1001861,

PLACE1001377, PLACE1001384, PLACE10013 87, PLACE1002493, PLACE1002591, PLACE1002685, PLACE1003353, PLACE1004128, PLACE1004 918, PLACE1005502, PLACE1005739, PLACE1005968, PLACE1006385, PLACE1007238, PLACE100 7375, PLACE1007488, PLACE1008244, PLACE1008650, PLACE1009319, PLACE1009468, PLACE10 10083, PLACE1010942, PLACE1011041, PLACE1011046, PLACE1011923, PLACE2000164, PLACE3 000145, PLACE3000350, PLACE3000475, THYRO1001595, Y79AA1000059, Y79AA1000966,

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(4) The following 230 clones were predicted to encode a protein in the category of proteins associated with transcription (including the clones belonging to plural categories). HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000851, 15 HEMBA1001137 , HEMBA1001510, HEMBA1001579, HEMBA1001800, HEMBA1001809, HEMBA1001819, HEMBA100184 7, HEMBA1002035, HEMBA1002092, HEMBA1002569, HEMBA1002609, HEMBA1003545, HEMBA10035 68, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004097, HEMBA1004321, HEMBA1004 353, HEMBA1004479, 20 HEMBA1004758, HEMBA1005359, HEMBA1005528, HEMBA1005548, HEMBA100 5931, HEMBA1006253, HEMBA1006278, HEMBA1006359, HEMBA1006559, HEMBB1000250, HEMBB10 00789, HEMBB1001011, HEMBB1001314, HEMBB1001482, HEMBB1001908, HEMBB1002134, HEMBB1 002217, MAMMA1000155, MAMMA1000183, MAMMA1000388, 25 MAMMA1001633, MAMMA1001820, MAMMA 1001837, MAMMA1002650, MAMMA1002937, NT2RM1000055, NT2RM1000666, NT2RM1000691, NT2R M1000894, NT2RM1001092, NT2RM2000013, NT2RM2000735, NT2RM2001035, NT2RM2001575, NT2 RM2001605, NT2RM2001670, NT2RM2001771, NT2RM2001797, NT2RM2002580, NT2RM4000024, NT 30 2RM4000104, NT2RM4000191, NT2RM4000496, NT2RM4000531, NT2RM4000734, NT2RM4000751, N T2RM4000996, NT2RM4001140, NT2RM4001200, NT2RM4001483, NT2RM4001592, NT2RM4001746, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RP1000111 , NT2RP1000333, NT2RP1000574, 35 NT2RP1000721, NT2RP1000860, NT2RP2000008, NT2RP200013 3, NT2RP2000297, NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP2002252, NT2RP20025 03, NT2RP2002591, NT2RP2003125, NT2RP2003228, NT2RP2003293, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2RP2004187, NT2RP2004961, NT2RP2005003, 40 NT2RP2005037 . NT2RP2005287, NT2RP2005496, NT2RP2005539, NT2RP2005722. NT2RP2006238, NT2RP200646 4, NT2RP3000050, NT2RP3000397, NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP30006 32, NT2RP3001057, NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001398, NT2RP3001 399, NT2RP3001671, 45 NT2RP3001672, NT2RP3001688, NT2RP3001712, NT2RP3001792, NT2RP300 1855, NT2RP3002165, NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193, NT2RP30 03251, NT2RP3003327, NT2RP3004258, NT2RP3004566, NT2RP3004572, NT2RP3004594. NT2RP3 004617. NT2RP4000078. NT2RP4000210. NT2RP4000398. 50 NT2RP4000449, NT2RP4000787, NT2RP 4000865, NT2RP4000997, NT2RP4001029, NT2RP4001213, NT2RP4001433, NT2RP4001529, NT2R P4001551, NT2RP4001638, NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP4002078, OVA RC1000479, OVARC1000846, OVARC1001381, OVARC1001555, OVARC1001577, OVARC1001949, PL 55 ACE1000066, PLACE1000133, PLACE1000583, PLACE1000596, PLACE1000706, PLACE1001118, P LACE1001238, PLACE1001304, PLACE1001602, PLACE1001632,

PLACE1002450, PLACE1002532, PLACE1002834, PLACE1003302, PLACE1003605,

PLACE1003738, PLACE1003885, PLACE1004471 , PLACE1004510, PLACE1004681, PLACE1006414, PLACE1006438, PLACE1006482, PLACE100723 9, PLACE1007346, PLACE1007544, PLACE1007598, PLACE1007955, PLACE1007969, PLACE10086 27, PLACE1008941, PLACE1009099, PLACE1009246, PLACE1009398, PLACE1010261, PLACE1010491, PLACE1010702, PLACE1010771, PLACE1010870, PLACE1011576, PLACE2000072 , PLACE2000359, PLACE4000128, PLACE4000156, PLACE4000259, PLACE4000431, THYRO100012 1, THYRO1000242, THYRO1000501, THYRO1000585, THYRO1001100, THYRO1001189, Y79AA10000 37, Y79AA1000627, Y79AA10010705, Y79AA1001105, Y79AA1001299, Y79AA1001391, Y79AA1001 533, Y79AA1001613, Y79AA1001848, Y79AA1001866, Y79AA1002093, Y79AA1002472, Y79AA100 2482.

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15 (5) The following 436 clones were predicted to encode a protein in the category of proteins associated with diseases (including the clones belonging to plural categories). HEMBA1000158, HEMBA1000216, HEMBA1000523, HEMBA1000561, HEMBA1000569, HEMBA1000673 , HEMBA1000972, HEMBA1001017, HEMBA1001019, HEMBA1001059, 20 HEMBA1001071, HEMBA100108 8, HEMBA1001123, HEMBA1001407, HEMBA1001569, HEMBA1001570, HEMBA1001672, HEMBA10020 35, HEMBA1002160, HEMBA1002162, HEMBA1002547, HEMBA1002939, HEMBA1003077, HEMBA1003 418, HEMBA1003433, HEMBA1003538, HEMBA1003555, HEMBA1003568, HEMBA1003569, HEMBA100 3581. 25 HEMBA1003720, HEMBA1003760, HEMBA1004168, HEMBA1004248, HEMBA1004276, HEMBA10 04354, HEMBA1004356, HEMBA1004479, HEMBA1004534, HEMBA1004669. HEMBA1004752, HEMBA1 004753, HEMBA1004807, HEMBA1005293, HEMBA1005331, HEMBA1005359, HEMBA1005367, HEMBA 1005423, HEMBA1005443, HEMBA1005475, 30 HEMBA1005528, HEMBA1005576, HEMBA1005581, HEMB A1005679, HEMBA1005699, HEMBA1005732, HEMBA1005931, HEMBA1006173, HEMBA1006474, HEM BA1006562, HEMBA1006648, HEMBA1006652, HEMBA1006737, HEMBA1006807, HEMBA1006976, HE MBA1007078, HEMBA1007121, HEMBA1007243, HEMBA1007301, HEMBB1000264. 35 HEMBB1000637, H EMBB1000684, HEMBB1000693, HEMBB1000915, HEMBB1000927, HEMBB1001011, HEMBB1001051, HEMBB1001068, HEMBB1001102, HEMBB1001119, HEMBB1001133, HEMBB1001177, HEMBB1001802, HEMBB1001908, HEMBB1001922, HEMBB1001967, HEMBB1002050, HEMBB1002193, HEMBB100221 7, HEMBB1002358, 40 HEMBB1002635, HEMBB1002683, MAMMA1000009, MAMMA1000092, MAMMA10001 03, MAMMA1000173, MAMMA1000731, MAMMA1000897, MAMMA1000956, MAMMA1001021. MAMMA1001041, MAMMA1001105, MAMMA1001198, MAMMA1001305, MAMMA1001576, MAMMA1001633, MAMMA1002140, MAMMA1002170, MAMMA1002198, MAMMA1002268, 45 MAMMA1002352, MAMMA100239 2, MAMMA1002485, MAMMA1002530, MAMMA1002573, MAMMA1002636, MAMMA1002858, MAMMA10028 69, MAMMA1002881, MAMMA1002937. MAMMA1003047, MAMMA1003099, MAMMA1003146, MAMMA1003 166, NT2RM1000035, NT2RM1000153, NT2RM1000257, NT2RM1000318, NT2RM1000377, NT2RM100 0394, 50 NT2RM1000555, NT2RM1000691, NT2RM1000725, NT2RM1000770, NT2RM1000826, NT2RM10 00850, NT2RM1001059, NT2RM1001072, NT2RM1001082, NT2RM1001092, NT2RM2000363, NT2RM2 000374, NT2RM2000594, NT2RM2000599, NT2RM2000714, NT2RM2000740, NT2RM2000821, NT2RM 2001035, NT2RM2001499, NT2RM2001575, 55 NT2RM2001605, NT2RM2001664, NT2RM2001670, NT2R M2001803, NT2RM2001823, NT2RM2001839, NT2RM4000085, NT2RM4000139, NT2RM4000155, NT2 RM4000290, NT2RM4000689, NT2RM4000895, NT2RM4001320, NT2RM4001566, NT2RM4001629, NT

2RM4001741, NT2RM4001819, NT2RM4001823, NT2RM4001865, NT2RM4001940, NT2RM4002093, N T2RM4002146, NT2RM4002161, NT2RM4002189, NT2RM4002457, NT2RM4002558, NT2RM4002571, NT2RP1000086, NT2RP1000376, NT2RP1000574, NT2RP1000581, NT2RP1000629, NT2RP1000721 , NT2RP1000733, NT2RP1000738, 5 NT2RP1000825, NT2RP1000833, NT2RP1000860, NT2RP100096 6, NT2RP1001033. NT2RP1001247, NT2RP2000126, NT2RP2000147, NT2RP2000233, NT2RP20002 97, NT2RP2000310, NT2RP2000414, NT2RP2000448, NT2RP2000459, NT2RP2000603. 10 NT2RP2000812, NT2RP2001127, NT2RP2001327, NT2RP2001440, NT2RP2001520. NT2RP2001663, NT2RP2001748, NT2RP2001876, NT2RP2001898, NT2RP2002025. NT2RP2002066, NT2RP200225 9, NT2RP2002312, NT2RP2002503, NT2RP2002618, NT2RP2003228, NT2RP2003293, NT2RP20032 95, NT2RP2003517, NT2RP2003564, NT2RP2003706, NT2RP2003737, NT2RP2003968, NT2RP2004 013, NT2RP2004098, NT2RP2004187, NT2RP2004316, NT2RP2004523, NT2RP2004538, NT2RP200 4933, NT2RP2005144, NT2RP2005288, NT2RP2005289, NT2RP2005325, NT2RP2005358, NT2RP20 05531, NT2RP2005773, NT2RP2006219, NT2RP3000050, NT2RP3000080, 20 NT2RP3000299, NT2RP3 000397, NT2RP3000512, NT2RP3000527, NT2RP3001081, NT2RP3001216. NT2RP3001268. NT2RP 3001355, NT2RP3001427, NT2RP3001428, NT2RP3001497, NT2RP3001554, NT2RP3001724, NT2R P3001764, NT2RP3001799, NT2RP3001855, NT2RP3002045, NT2RP3002151, NT2RP3002351, NT2 RP3002352, 25 NT2RP3002663, NT2RP3002818, NT2RP3002876, NT2RP3002909, NT2RP3002972, NT 2RP3003061, NT2RP3003157, NT2RP3003242, NT2RP3003282, NT2RP3003672. NT2RP3004078, N T2RP3004209, NT2RP3004399, NT2RP3004424, NT2RP3004475, NT2RP3004480, NT2RP3004490, NT2RP3004527, NT2RP3004578, NT2RP4000049, 30 NT2RP4000109, NT2RP4000185, NT2RP4000312 , NT2RP4000367, NT2RP4000398, NT2RP4000457, NT2RP4000855, NT2RP4000879, NT2RP400091 5, NT2RP4000928, NT2RP4001095, NT2RP4001150, NT2RP4001213, NT2RP4001483, NT2RP40014 98, NT2RP4001753, NT2RP4001838, OVARC1000014, OVARC1000114, OVARC1000139. 35 OVARC1000241, OVARC1000440, OVARC1000543, OVARC1000722, OVARC1000771. OVARC1000834 , OVARC1000850, OVARC1001113, OVARC1001244, OVARC1001296, OVARC1001357, OVARC100137 2, OVARC1001417, OVARC1001496, OVARC1001506, OVARC1001577, OVARC1001668, OVARC10017 13, OVARC1001809, OVARC1001880, 40 OVARC1001901, OVARC1001949, OVARC1002050, PLACE1000 133, PLACE1000142, PLACE1000184, PLACE1000383, PLACE1000406, PLACE1000420, PLACE100 0583, PLACE1000588, PLACE1000706, PLACE1001384, PLACE1001387, PLACE1001517, PLACE10 01602, PLACE1001634, PLACE1001771, PLACE1002046, PLACE1002437, 45 PLACE1002474, PLACE1 002665, PLACE1002834, PLACE1003302, PLACE1003493, PLACE1003704, PLACE1003888, PLACE 1003903, PLACE1003968, PLACE1004197, PLACE1004358, PLACE1004471, PLACE1004681, PLAC E1004751, PLACE1004773, PLACE1005101, PLACE1006040, PLACE1006119, PLACE1006170, PLA CE1006187, . 50 PLACE1006288, PLACE1006414, PLACE1006438, PLACE1006962, PLACE1007239, PL ACE1007257, PLACE1007488, PLACE1007511, PLACE1007737, PLACE1008000, PLACE1008402, P LACE1008643, PLACE1008696, PLACE1008941, PLACE1009027, PLACE1009200, PLACE1009230, PLACE1009298, PLACE1009319, PLACE1009444, 55 PLACE1009524, PLACE1010083, PLACE1010089, PLACE1010599, PLACE1010833, PLACE1010856, PLACE1011054, PLACE1011057, PLACE101116 5, PLACE1011229, PLACE1011263, PLACE1011371, PLACE2000072, PLACE2000399, PLACE20004 38,

PLACE2000458, PLACE3000004, PLACE3000242, PLACE3000352, PLACE3000353, PLACE3000455, PLACE3000475, PLACE4000009, PLACE4000014, PLACE4000052, PLACE4000320, PLACE4000581, SKNMC1000050, THYRO1000026, THYRO1000173, THYRO1000186, THYRO100032 0, THYRO1000327, THYRO1000387, THYRO1000471, THYRO1000934, THYRO1001458, THYRO10015 95, THYRO1001605, THYRO1001617, THYRO1001671, Y79AA1000037, Y79AA1000469, Y79AA1000 539, Y79AA1000560, Y79AA1000734, Y79AA1000985, Y79AA1001105, Y79AA1001391, Y79AA100 1394, Y79AA1001548, Y79AA1001613, Y79AA1001874, Y79AA1002258, Y79AA1002416, Y79AA10 02482,

(6) The following 301 clones were predicted to encode a protein in the category of enzyme or 15 proteins associated with metabolism (including the clones belonging to plural categories). HEMBA1000012. HEMBA1000459. HEMBA1000852. HEMBA1001059. HEMBA1001257. HEMBA1001620 , HEMBA1001714, HEMBA1001744, HEMBA1001896, HEMBA1002003, HEMBA1002212, HEMBA100225 7, HEMBA1002513, HEMBA1002973, HEMBA1003046, 20 HEMBA1003136, HEMBA1003179, HEMBA10032 86, HEMBA1003291, HEMBA1003538, HEMBA1003680, HEMBA1004227, HEMBA1004408, HEMBA1004 734, HEMBA1004795, HEMBA1005513, HEMBA1005737, HEMBA1005815, HEMBA1006278, HEMBA100 6347, HEMBA1006521, HEMBA1006648, HEMBA1006877, HEMBA1006941, HEMBA1006976, 25 HEMBA10 07121, HEMBA1007243, HEMBA1007300, HEMBB1000781, HEMBB1000915, HEMBB1001288, HEMBB1 001429, HEMBB1001443, HEMBB1001915, HEMBB1002042, HEMBB1002342, HEMBB1002358, HEMBB 1002635, HEMBB1002683, MAMMA1000020, MAMMA1000085, MAMMA1000672, MAMMA1000841, MAMM A1001008, MAMMA1001139, 30 MAMMA1001476, MAMMA1001501, MAMMA1002268, MAMMA1002530, MAM MA1002619, MAMMA1002671, MAMMA1002938, NT2RM1000132, NT2RM1000153, NT2RM1000256, NT 2RM1000280, NT2RM1000377, NT2RM1000648, NT2RM1000850, NT2RM1000867, NT2RM1000894, N T2RM1001072, NT2RM2000013, NT2RM2000124, NT2RM2000191, 35 NT2RM2000368, NT2RM2000371, NT2RM2000577, NT2RM2000594, NT2RM2000599, NT2RM2000951, NT2RM2001238, NT2RM2001664, NT2RM2001700, NT2RM2001730, NT2RM2001785, NT2RM2001803, NT2RM2002030, NT2RM400002 4, NT2RM4000155, NT2RM4000344, NT2RM4000496, NT2RM4000616, NT2RM4000712, NT2RM40008 95, 40 NT2RM4001313, NT2RM4001316, NT2RM4001340, NT2RM4001444, NT2RM4001758, NT2RM4001819, NT2RM4001930, NT2RM4002062, NT2RM4002063, NT2RM4002409, NT2RM4002558 , NT2RM4002571, NT2RM4002623, NT2RP1000018, NT2RP1000112, NT2RP1000376, NT2RP100060 9, NT2RP1000833, NT2RP1000834, NT2RP1001079, 45 NT2RP1001253, NT2RP1001361, NT2RP10015 43, NT2RP2000056, NT2RP2000114, NT2RP2000183, NT2RP2000329, NT2RP2000422, NT2RP2000 710, NT2RP2000764, NT2RP2000932, NT2RP2001070, NT2RP2001663, NT2RP2001839, NT2RP200 1898, NT2RP2002256, NT2RP2002312, NT2RP2002442, NT2RP2002595, NT2RP2002618, 50 NT2RP20 02993, NT2RP2003230, NT2RP2003286, NT2RP2003506, NT2RP2003643, NT2RP2003706, NT2RP2 003737, NT2RP2003781, NT2RP2003912, NT2RP2004232, NT2RP2004239, NT2RP2004768, NT2RP 2004791, NT2RP2004799, NT2RP2005038, NT2RP2005276, NT2RP2005344, NT2RP2005457, NT2R P2005498, NT2RP2005531, 55 NT2RP2005557, NT2RP2005690, NT2RP2005773, NT2RP2005775, NT2 RP2005942, NT2RP2006571, NT2RP3000046, NT2RP3000085, NT2RP3000359, NT2RP3000418, NT

2RP3000742, NT2RP3000845, NT2RP3000875, NT2RP3001274, NT2RP3001407,

NT2RP3001495, N T2RP3002351, NT2RP3002402, NT2RP3002969, NT2RP3003301, NT2RP3003346, NT2RP3003500, NT2RP3003800, NT2RP3003825, NT2RP3003831, NT2RP3003914, NT2RP3004155, NT2RP3004209 NT2RP3004669, NT2RP3004670, 5 NT2RP4000259, NT2RP4000417, NT2RP4000457, NT2RP400072 4, NT2RP4000855, NT2RP4000928, NT2RP4000997, NT2RP4001041, NT2RP4001079, NT2RP40010 95, NT2RP4001219, NT2RP4001345, NT2RP4001375, NT2RP4001483, NT2RP4001592, NT2RP4001644. NT2RP4001893. NT2RP4001946. NT2RP4002408. NT2RP5003500. 10 NT2RP5003522 , OVARC100060, OVARC1000139, OVARC1000309, OVARC1000543, OVARC1000682, OVARC100072 2, OVARC1000885, OVARC1001261, OVARC1001610, OVARC1001713, OVARC1001762, OVARC10018 09, OVARC1001942, PLACE1000007, PLACE1000142, PLACE1000420, PLACE1000547, PLACE1000 653, PLACE1001062, 15 PLACE1001603, PLACE1001692, PLACE1001817, PLACE1001989, PLACE100 2991, PLACE1003100, PLACE1003709, PLACE1003885, PLACE1003888, PLACE1003903, PLACE10 03915, PLACE1003968, PLACE1004428, PLACE1004437, PLACE1004751, PLACE1005305, PLACE1 005477, PLACE1005656, PLACE1005763, PLACE1005804, 20 PLACE1006011, PLACE1016469, PLACE 1006731, PLACE1007729, PLACE1007958, PLACE1008275, PLACE1008356, PLACE1008696, PLAC E1009444, PLACE1009861, PLACE1009997, PLACE1010089, PLACE1010096, PLACE1010106, PLA CE1010152, PLACE1010481, PLACE1011046, PLACE1011203, PLACE1011219, PLACE1011229, PL 25 ACE1011332, PLACE1011923, PLACE2000021, PLACE2000140, PLACE2000404. PLACE2000411, P LACE3000020, PLACE3000148, PLACE3000156, PLACE3000350, PLACE3000353, PLACE4000259, PLACE4000431, PLACE4000558, PLACE4000590, SKNMC1000050, THYRO1000017, THYRO1000197, THYRO1000320, THYRO1000358, 30 THYRO1000471, THYRO1000926, THYRO1000934, THYRO100095 1, THYRO1001406, THYRO1001617, THYRO1001671, Y79AA1001048, Y79AA1001233, Y79AA10015 81, Y79AA1001679, Y79AA1001827, Y79AA1002027, Y79AA1002209, Y79AA1002361, Y79AA1002 416,

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(7) The following 74 clones were predicted to encode a protein in the category of proteins associated with cell division or cell proliferation (including the clones belonging to plural categories).

HEMBA1001019, HEMBA1001595, HEMBA1002229, HEMBA1002363, HEMBA1003136, HEMBA1003836 , HEMBA1004131, HEMBA1005241, HEMBB1000399, HEMBB1001175, HEMBB1001242, MAMMA100182 0, MAMMA1003011, NT2RM1000394, NT2RM2001256, NT2RM4000215, NT2RM4001611, NT2RM40017 14, NT2RP1000733, NT2RP1001177, NT2RP2000077, NT2RP2000346, NT2RP2001397, NT2RP2003 912, NT2RP2005520, NT2RP2005675, NT2RP2005767, NT2RP2005857, NT2RP2006464, NT2RP300 0031, NT2RP3001155, NT2RP3001730, NT2RP3002151, NT2RP3002273, NT2RP3002399, NT2RP30 04507, NT2RP3004578, NT2RP3004594, NT2RP4000210, NT2RP4000498, NT2RP4001207, NT2RP4 001414, NT2RP4001551, OVARC1000006, OVARC1000087, OVARC1000113, OVARC1000846, OVARC 1000876, OVARC1001068, OVARC1001107, OVARC1001113, OVARC1002112, PLACE1000547, PLAC E1001611, PLACE1001691, PLACE1002775, PLACE1002811, PLACE1003709, PLACE1004316, PLA CE1004674.

PLACE1004814, PLACE1005287, PLACE1006187, PLACE1006368, PLACE1008808, PLACE1008947, PLACE1009060, PLACE1010720, PLACE1010761, PLACE1010833, PLACE1011056, PLACE3000244, Y79AA1000214, Y79AA1001236,

- (8) The following 92 clones were predicted to encode a protein in the category of proteins associated with cytoskeleton (including the clones belonging to plural categories).
- HEMBA1000020, HEMBA1000588, HEMBA1001804, HEMBA1002102, HEMBA1002161, HEMBA1002939 , HEMBA1003235, HEMBA1003581, HEMBA1004534, HEMBA1005009, HEMBA1005595, HEMBA100627 2, HEMBA1006737, HEMBA1007018, HEMBB1000404, HEMBB1001282, MAMMA1000173, MAMMA10004 29, MAMMA1001041, MAMMA1001576, MAMMA1001735, MAMMA1002622, MAMMA1002637, MAMMA1003 099, MAMMA1003127, NT2RM1000898, NT2RM1001003, NT2RM1001139, NT2RM2000691, NT2RM400 0030, NT2RM4000167, NT2RM4000689, NT2RM4001741, NT2RM4001776, NT2RM4002109, NT2RP10 00348, NT2RP1000460, NT2RP1000478, NT2RP1001033, NT2RP2000812,
- 15 NT2RP2001634, NT2RP2 001900, NT2RP2002727, NT2RP2003307, NT2RP2003604, NT2RP2004538, NT2RP2005531, NT2RP 3000868, NT2RP3001216, NT2RP3001272, NT2RP3002688, NT2RP3003061, NT2RP3003138, NT2R P3003230, NT2RP3003282,
- NT2RP3004454, NT2RP3004569, NT2RP4001126, NT2RP4001927, OVA RC1000437, OVARC1000520, OVARC1000679, OVARC1001731, OVARC1002050, PLACE1002571, PL ACE1002591, PLACE1002655, PLACE1003669, PLACE1004777, PLACE1005287, PLACE1006368, P LACE1007105, PLACE1010310, PLACE1010743, PLACE1010960, PLACE1011263, PLACE2000039, PLACE2000216, PLACE2000274, PLACE3000145.
- PLACE3000322, PLACE4000009, PLACE4000612, THYRO1000072, THYRO1000666, THYRO1001121, THYRO1001458, Y79AA1000368, Y79AA100083 3, Y79AA1000962, Y79AA1000985, Y79AA1002208.
- (9) The following 280 clones were predicted to encode a protein in the category of proteins associated with RNA synthesis (including the clones belonging to plural categories). HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000591,
- HEMBA1000851 , HEMBA1001137, HEMBA1001510, HEMBA1001579, HEMBA1001800, HEMBA1001809, HEMBA100181 9, HEMBA1001847, HEMBA1002035, HEMBA1002092, HEMBA1002569, HEMBA1002609, HEMBA10034 18, HEMBA1003545, HEMBA1003568, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004 097, HEMBA1004321,
- HEMBA1004353, HEMBA1004479, HEMBA1004669, HEMBA1004758, HEMBA100 5359, HEMBA1005528, HEMBA1005548, HEMBA1005931, HEMBA1006253, HEMBA1006278,
- HEMBA1003226, HEMBA1003348, HEMBA1003931, HEMBA1000233, HEMBA1000278, HEMBA10 06359, HEMBA1006559, HEMBA1006639, HEMBB1000250, HEMBB1000789, HEMBB1001011, HEMBB1 001314, HEMBB1001482, HEMBB1001908, HEMBB1002134, HEMBB1002217, MAMMA1000155, MAMMA 1000183, MAMMA1000388, MAMMA1001059,
- MAMMA1001633, MAMMA1001820, MAMMA1001837, MAMM A1002650, MAMMA1002937, NT2RM1000055, NT2RM1000187, NT2RM1000666, NT2RM1000691, NT2 RM1000852, NT2RM1000894, NT2RM1001092, NT2RM200013, NT2RM2000735, NT2RM2001035, NT2RM2001424, NT2RM2001575, NT2RM2001605, NT2RM2001670, NT2RM2001771,
- NT2RM2001797, N T2RM2001989, NT2RM2002100, NT2RM2002580, NT2RM4000024, NT2RM4000104, NT2RM4000191, NT2RM4000421, NT2RM4000471, NT2RM4000496, NT2RM4000531, NT2RM4000734, NT2RM4000751 , NT2RM4000996, NT2RM4001140, NT2RM4001200, NT2RM4001483, NT2RM4001592, NT2RM400174 6, NT2RM4001783.
- 55 NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RM40020 93, NT2RM4002479, NT2RP1000111, NT2RP1000272, NT2RP1000333, NT2RP1000470, NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP100180, NT2RP2000008.

NT2RP2000133 , NT2RP2000161, NT2RP2000297, NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP200144 9, NT2RP2002099, NT2RP2002252, NT2RP2002503, NT2RP2002591, NT2RP2002928, NT2RP20031 25, NT2RP2003228, NT2RP2003293. NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2RP2004 187, NT2RP2004568, 5 NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005126, NT2RP200 5168, NT2RP2005239, NT2RP2005287, NT2RP2005496, NT2RP2005539, NT2RP2005722, NT2RP20 05763, NT2RP2005942, NT2RP2006238, NT2RP2006464, NT2RP3000050. 10 NT2RP3000361, NT2RP3 000397, NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP3000632, NT2RP3000917, NT2RP 3001057, NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001398, NT2RP3001399, NT2R P3001671, NT2RP3001672, NT2RP3001688, NT2RP3001712, NT2RP3001792, NT2RP3001855, NT2 RP3002165, 15 NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193, NT2RP3003251, NT 2RP3003327, NT2RP3004013, NT2RP3004258, NT2RP3004504, NT2RP3004566, NT2RP3004572, N T2RP3004594, NT2RP3004617, NT2RP4000078, NT2RP4000111, NT2RP4000210, NT2RP4000398, NT2RP4000449, NT2RP4000518, NT2RP4000614, NT2RP4000787, NT2RP4000865, NT2RP4000997 , NT2RP4001029, NT2RP4001080, 20 NT2RP4001148, NT2RP4001213, NT2RP4001433, NT2RP400152 9, NT2RP4001551, NT2RP4001638, NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP40020 78, OVARC1000479, OVARC1000846, OVARC1001381, OVARC1001555, OVARC1001577, 25 OVARC1001949, PLACE1000066, PLACE1000133, PLACE1000583, PLACE1000596, PLACE1000706 , PLACE1001118, PLACE1001238, PLACE1001304, PLACE1001602, PLACE1001632, PLACE100245 0, PLACE1002532, PLACE1002834, PLACE1003190, PLACE1003302, PLACE1003605, PLACE10037 04, PLACE1003738, PLACE1003885, 30 PLACE1004471, PLACE1004510, PLACE1004564, PLACE1004 681, PLACE1004902, PLACE1005646, PLACE1005876, PLACE1006196, PLACE1006414, PLACE100 6438, PLACE1006482, PLACE1007239, PLACE1007346, PLACE1007544, PLACE1007598. PLACE10 07955, PLACE1007969, PLACE1008627, PLACE1008941, PLACE1009099, 35 PLACE1009246, PLACE1 009398, PLACE1010053, PLACE1010194, PLACE1010261, PLACE1010491, PLACE1010580, PLACE 1010702, PLACE1010771, PLACE1010870, PLACE1011576, PLACE2000072, PLACE2000359, PLAC E4000128, PLACE4000156, PLACE4000259, PLACE4000431, THYRO1000121, THYRO1000242, THY RO1000501, 40 THYRO1000585, THYRO1001100, THYRO1001189, Y79AA1000037, Y79AA1000349, Y7 Y79AA1000627, Y79AA1000705, Y79AA1000752, 9AA1000539. Y79AA1001105. Y79AA1001299, Y 79AA1001391, Y79AA1001533, Y79AA1001613, Y79AA1001848, Y79AA1001866, Y79AA1001963, Y79AA1002093, Y79AA1002472, Y79AA1002482,

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(10) The following 352 clones were predicted to encode a protein in the category of nuclear protein (including the clones belonging to plural categories).

HEMBA100005, HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000827, HEMBA1000851, HEMBA1001088, HEMBA1001137, HEMBA1001476, HEMBA1001510, HEMBA100157, HEMBA1001800, HEMBA1001809, HEMBA1001819, HEMBA1001847, HEMBA1002035, HEMBA10020, 92, HEMBA1002363, HEMBA1002495, HEMBA1002569, HEMBA1002609, HEMBA1002999, HEMBA1003, 148, HEMBA1003418, HEMBA1003545, HEMBA1003568, HEMBA1003684, HEMBA1003760, HEMBA100, 3783, HEMBA1003953, HEMBA1004097, HEMBA1004321, HEMBA1004353, HEMBA1004479, HEMBA10, 04669, HEMBA1004758, HEMBA1005559, HEMBA1005575, HEMBA1005528,

HEMBA1005548, HEMBA1 005931, HEMBA1006253, HEMBA1006278, HEMBA1006359, HEMBA1006474, HEMBA1006559, HEMBA 1006914, HEMBB1000250, HEMBB1000399, HEMBB1000789, HEMBB1001011, HEMBB1001314, HEMB B1001482, HEMBB1001908, HEMBB1001915, HEMBB1002134, HEMBB1002217, MAMMA1000155, MAM MA1000183. MAMMA1000388, MAMMA1000731, MAMMA1001633, MAMMA1001820, MAMMA1001837, MA MMA1002650, MAMMA1002869, MAMMA1002937, MAMMA1003011, NT2RM1000055. NT2RM1000187, N T2RM1000257, NT2RM1000394, NT2RM1000666, NT2RM1000691, NT2RM1000852, NT2RM1000894, NT2RM1001092, NT2RM2000013, NT2RM2000735. 10 NT2RM2001035, NT2RM2001575, NT2RM2001605 , NT2RM2001635, NT2RM2001670, NT2RM2001771, NT2RM2001797, NT2RM2001823, NT2RM200193 6, NT2RM2001989, NT2RM2002100, NT2RM2002580, NT2RM4000024, NT2RM4000085, NT2RM40001 04, 15 NT2RM4000191, NT2RM4000215, NT2RM4000290, NT2RM4000421, NT2RM4000496, NT2RM4000531, NT2RM4000734, NT2RM4000751, NT2RM4000996, NT2RM4001140. NT2RM4001200 , NT2RM4001217, NT2RM4001382, NT2RM4001483, NT2RM4001592, NT2RM4001605, NT2RM400161 1, NT2RM4001746, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM40019 79, NT2RM4002093, NT2RM4002146, 20 NT2RM4002527, NT2RP1000035, NT2RP1000111, NT2RP1000 333, NT2RP1000493, NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP1000966, NT2RP100 1177, NT2RP2000008, NT2RP2000126, NT2RP2000133, NT2RP2000161, NT2RP2000297, 25 NT2RP20 00414, NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP2001420, NT2RP2001449, NT2RP2 001536, NT2RP2001762, NT2RP2002185, NT2RP2002252, NT2RP2002259, NT2RP2002503, NT2RP 2002504, NT2RP2002591, NT2RP2002993, NT2RP2003125, NT2RP2003228, NT2RP2003277, NT2R P2003286, NT2RP2003293, 30 NT2RP2003308, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2 RP2004187. NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005287, NT2RP2005496, NT 2RP2005520, NT2RP2005539, NT2RP2005722, NT2RP2005767. NT2RP2005857. NT2RP2005942, N T2RP2006238, NT2RP2006464, NT2RP3000031, NT2RP3000050, 35 NT2RP3000361, NT2RP3000397, NT2RP3000512, NT2RP3000527, NT2RP1000590. NT2RP3000603, NT2RP3000632, NT2RP3000917 , NT2RP3001057, NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001274, NT2RP300139 8, NT2RP3001399, NT2RP3001427, NT2RP3001428, NT2RP3001587, NT2RP3001671, NT2RP30016 72, 40 NT2RP3001688, NT2RP3001712, NT2RP3001724, NT2RP3001792, NT2RP3001855, NT2RP3002056, NT2RP3002165, NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193 , NT2RP3003212, NT2RP3003251, NT2RP3003327, NT2RP3004206, NT2RP3004209, NT2RP300425 8, NT2RP3004566, NT2RP3004572, NT2RP3004594, 45 NT2RP3004617, NT2RP4000078, NT2RP40001 11, NT2RP4000150, NT2RP4000210, NT2RP4000398, NT2RP4000449, NT2RP4000457, NT2RP4000 518, NT2RP4000787, NT2RP4000865, NT2RP4000997, NT2RP4001029, NT2RP4001080, NT2RP400 1148, NT2RP4001207, NT2RP4001213, NT2RP4001433, NT2RP4001529, NT2RP4001551, 50 NT2RP40 01638, NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP4002078, OVARC1000006, OVARC1 000087, OVARC1000139, OVARC1000326, OVARC1000440, OVARC1000479, OVARC1000846, OVARC 1000883, OVARC1001180, OVARC1001244, OVARC1001381, OVARC1001555, OVARC1001577, OVAR C1001702, OVARC1001949. 55 OVARC1002112, PLACE1000066, PLACE1000133, PLACE1000406, PLA CE1000583, PLACE1000596, PLACE1000706, PLACE1001118, PLACE1001238, PLACE1001304, PL ACE1001602, PLACE1001611, PLACE1001632, PLACE1002450, PLACE1002532.

PLACE1002834, P LACE1003100, PLACE1003190, PLACE1003302, PLACE1003605, PLACE1003704, PLACE1003738, PLACE1003885, PLACE1004471, PLACE1004510, PLACE1004564, PLACE1004681, PLACE1004902, PLACE1005287, PLACE1005876, PLACE1006011, PLACE1006119, PLACE1006414, PLACE100643 8, PLACE1006482, 5 PLACE1007239, PLACE1007346, PLACE1007544, PLACE1007598, PLACE10079 55, PLACE1007969, PLACE1008044, PLACE1008177, PLACE1008603, PLACE1008627, PLACE1008941, PLACE1008947, PLACE1009099, PLACE1009113, PLACE1009246, 10 PLACE1009398 , PLACE1010152, PLACE1010194, PLACE1010261, PLACE1010491, PLACE1010702, PLACE101072 0, PLACE1010771, PLACE1010870, PLACE1011056, PLACE1011229, PLACE1011576, PLACE10116 64, PLACE2000072, PLACE2000359, PLACE2000411, PLACE4000014, PLACE4000128, PLACE4000 156, PLACE4000259, PLACE4000261, PLACE4000431, THYRO1000121, THYRO1000242, THYRO100 0501, 15 THYRO1000585, THYRO1001100, THYRO1001189, Y79AA1000037, Y79AA1000214, Y79AA10 00539, Y79AA1000589, Y79AA1000627, Y79AA1000705, Y79AA1000752, Y79AA1000784, Y79AA1 001105, Y79AA1001299, Y79AA1001391, Y79AA1001533, Y79AA1001613, Y79AA1001848, Y79AA 1001866, Y79AA1001963, Y79AA1002093, 20 Y79AA1002472, Y79AA1002482,

associated with protein synthesis or transport (including the clones belonging to plural categories).
 HEMBA1000012, HEMBA1000542, HEMBA1001913, HEMBA1003773, HEMBA1004202, HEMBA1004356
 HEMBA1004847, HEMBA1005047,HEMBA1005202, HEMBA1005963, HEMBA1006310, HEMBA100665
 JEMBA1006914, HEMBB1000217, HEMBB1000725,

(11) The following 111 clones were predicted to encode a protein in the category of proteins

- HEMBA1006310, HEMBA100665 2, HEMBA1006914, HEMBB1000217, HEMBB1000725, HEMBB1001346, HEMBB1001831, MAMMA10000 85, MAMMA1000734, MAMMA1002170, MAMMA1002236, MAMMA1002598, NT2RM1000661, NT2RM2000 504, NT2RM2000577, NT2RM2001131, NT2RM4000155, NT2RM4001444, NT2RM4002062, NT2RM400 2205, NT2RM4002623, NT2RP1000326, NT2RP1000947, NT2RP1001569, NT2RP2000710,
- NT2RP20 00880, NT2RP2001290, NT2RP2001511, NT2RP2001660, NT2RP2002185, NT2RP2002606, NT2RP2 002959, NT2RP2002980, NT2RP2003158, NT2RP2003760, NT2RP2004194, NT2RP2004791, NT2RP 2005012, NT2RP2005116, NT2RP2005204, NT2RP2006565, NT2RP2006598, NT2RP3000047, NT2R P3000366, NT2RP3000759, NT2RP3000968, NT2RP3001587, NT2RP3002529, NT2RP3003385, NT2 RP3003589, NT2RP3003876, NT2RP3004618, NT2RP4000370, NT2RP4000524, NT2RP4000528, NT
- 2RP4001041, NT2RP4001574, NT2RP4001592, NT2RP4002047, OVARC1000004, OVARC1000771, O VARC1000862, OVARC1001180, OVARC1001342, OVARC1001766, OVARC1002138, PLACE1000061, PLACE1000492, PLACE1000863, PLACE1001748, PLACE1002090, PLACE1003394, PLACE1003915 , PLACE1004149, PLACE1004743, PLACE1005557, PLACE1005813, PLACE1006488, PLACE1006653 1, PLACE1006615,
- PLACE1007706, PLACE1008273, PLACE1008790, PLACE1008813, PLACE10097 21, PLACE1009763, PLACE1009845, PLACE1010074, PLACE1011109, PLACE1011477, PLACE2000404, PLACE3000059, PLACE3000121, PLACE4000558, PLACE4000654, SKNMC1000011, THYRO1000288, THYRO1000974, Y79AA1000346, Y79AA1000968,

⁵⁵ Y79AA1002209,

(12) The following 23 clones were predicted to encode a protein in the category of proteins

associated with cellular defense (including the clones belonging to plural categories).

HEMBA100005, HEMBA1000531, HEMBB1000217, HEMBB1000399, MAMMA1000734, NT2RP1000493, NT2RP2000006, NT2RP2000045, NT2RP2001536, NT2RP2005204, NT2RP300059 0, NT2RP3001426, NT2RP3002056, NT2RP3004262, NT2RP4001638, OVARC1001900, PLACE10056 11, PLACE1006958, PLACE1008275, PLACE1009113, PLACE4000014, Y79AA1002139,

(13) The following 23 clones were predicted to encode a protein in the category of proteins associated with development or growth (including the clones belonging to plural categories).
 HEMBB1001802, HEMBB1002442, NT2RM2002142, NT2RM4001047, NT2RP2003308, NT2RP2004816, NT2RP3000994, NT2RP3001340, NT2RP3004206, NT2RP3004472,
 NT2RP4000246, NT2RP400100 6, OVARC1000304, OVARC1000746, OVARC1000996, PLACE1006037, PLACE1009622, PLACE10116 64, Y79AA1001692,

Table 9
Selected clones having the maximal ATGpr1 score of 0.3 or higher (3690 clones).

	clone	name of	maximal
	name	sequence	ATGpr1
5			score
	HEMBA1000005	F-HEMBA1000005	0. 84
	HEMBA1000012	F-HEMBA1000012	0. 56
	HEMBA1000020	F-HEMBA1000020	0. 94
	HEMBA1000030	F-HEMBA1000030	0. 44
	HEMBA1000046	F-HEMBA1000046	0.50
	HEMBA1000050	F-HEMBA1000050	0.94
	HEMBA1000076	F-HEMBA1000076	0.48
	HEMBA1000129	F-HEMBA1000129	0.74
	HEMBA1000141	F-HEMBA1000141	0.55
9	HEMBA1000150	F-HEMBA1000150	0. 72
	HEMBA1000156	F-HEMBA1000156	0. 94
	HEMBA1000158	F-HEMBA1000158	0. 62
	HEMBA1000168	F-HEMBA1000168	0. 94
5	HEMBA1000185	F-HEMBA1000185	0. 86
	HEMBA1000193	F-HEMBA1000193	0. 94

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		1 07 4 017 AZ	
	HEMBA1000201	F-HEMBA1000201	0.83
	HEMBA1000216	F-HEMBA1000216	0.32
	HEMBA1000227	F-HEMBA1000227	0.72
5	HEMBA1000231	F-HEMBA1000231_	0, 34
	HEMBA1000244	F-HEMBA1000244	0.51
	HEMBA1000288	F-HEMBA1000288	0.36
	HEMBA1000290	F-HEMBA1000290	0. 45
10	HEMBA1000302	F-HEMBA1000302	0.71
	HEMBA1000303	F-HEMBA1000303	0. 42
	HEMBA1000304	F-HEMBA1000304	0. 45
	HEMBA1000307	F-HEMBA1000307	0.48
15	HEMBA1000327	F-HEMBA1000327	0.94
	HEMBA1000356	F-HEMBA1000356	0.90
	HEMBA1000369	F-HEMBA1000369,	0. 94
	HEMBA1000387	F-HEMBA1000387	0.66
20	HEMBA1000392	F-HEMBA1000392	0.60
20	HEMBA1000396	F-HEMBA1000396	0. 81
	HEMBA1000456	F-HEMBA1000456	0.90
	HEMBA1000460	F-HEMBA1000460	0.31
05	HEMBA1000488	F-HEMBA1000488	0.47
25	HEMBA1000490	F-HEMBA1000490	0.89
	HEMBA1000491	F-HEMBA1000491	0.94
	HEMBA1000501	F-HEMBA1000501	0.67
	HEMBA1000505	F-HEMBA1000505	0.65
30	HEMBA1000508	F-HEMBA1000508	0.39
	HEMBA1000520	F-HEMBA1000520	0.94
	HEMBA1000523	F-HEMBA1000523	0.86
	HEMBA1000531	F-HEMBA1000531	0.49
35	HEMBA1000534	F-HEMBA1000534	0.52
	HEMBA1000542	F-HEMBA1000542	0.81
	HEMBA1000555	F-HEMBA1000555	0.67
	HEMBA1000561	F-HEMBA1000561	0.34
40	HEMBA1000568	F-HEMBA1000568	0.66
	HEMBA1000588	F-HEMBA1000588	0.79
	HEMBA1000591	F-HEMBA1000591	0.92
	HEMBA1000592	F-HEMBA1000592	0.73
45	HEMBA1000594	F-HEMBA1000594	0.89
	HEMBA1000608	F-HEMBA1000608	0.94
		F-HEMBA1000636	0.94
	HEMBA1000637		0. 92
50	HEMBA1000657	F-HEMBA1000657	0.85
	HEMBA1000682		0.94
	HEMBA1000686		0.90
	HEMBA1000719		0.94
55	HEMBA1000727	F-HEMBA1000727	0.40
	HEMBA1000752	F-HEMBA1000752	0. 31

HEMBA1000817 F-HEMBA1000817 0.94 HEMBA1000851 HEMBA1000851 0.63 HEMBA1000869 F-HEMBA1000869 0.82 HEMBA1000872 F-HEMBA1000872 0.54 HEMBA1000910 F-HEMBA1000910 0.91 HEMBA1000911 F-HEMBA1000911 0.83 HEMBA1000942 F-HEMBA1000912 0.31 HEMBA1000945 F-HEMBA1000919 0.83 HEMBA1000946 F-HEMBA1000968 0.75 HEMBA1000966 F-HEMBA1000971 0.74 HEMBA1000971 F-HEMBA1000975 0.46 HEMBA1000971 F-HEMBA1000975 0.46 HEMBA1000975 F-HEMBA1000975 0.46 HEMBA1001008 F-HEMBA1000975 0.46 HEMBA1001008 F-HEMBA1001008 0.47 HEMBA1001008 F-HEMBA1001009 0.85 HEMBA1001009 F-HEMBA1001009 0.85 HEMBA1001009 F-HEMBA1001009 0.86 HEMBA1001050 F-HEMBA1001050 0.77 HEMBA1001052 F-HEMBA1001052 0.83 HEMBA1001053 F-HEMBA1001055 0.78 HEMBA1001109 F-HEMBA1001050 0.77 HEMBA1001109 F-HEMBA1001050 0.78 HEMBA1001109 F-HEMBA1001050 0.78 HEMBA1001109 F-HEMBA1001050 0.79 HEMBA1001109 F-HEMBA1001050 0.79 HEMBA1001109 F-HEMBA1001050 0.70 HEMBA1001109 F-HEMBA1001050 0.50 HEMBA1001109 F-HEMBA1001050 0.50 HEMBA1001109 F-HEMBA1001133 0.45 HEMBA1001121 F-HEMBA1001133 0.45 HEMBA1001122 F-HEMBA1001133 0.45 HEMBA1001124 F-HEMBA1001130 0.65 HEMBA1001125 F-HEMBA1001130 0.65 HEMBA1001247 F-HEMBA1001257 0.92 HEMBA1001247 F-HEMBA1001250 0.94 HEMBA1001300 F-HEMBA1001300 0.79 HEMBA1001301 F-HEMBA1001301 0.66 HEMBA1001302 F-HEMBA1001303 0.79 HEMBA1001303 F-HEMBA1001307 0.94 HEMBA1001303 F-HEMBA1001307 0.94				
HEMBA1000851		HEMBA1000817	F-HEMBA1000817	0. 94
HEMBA1000869		HEMBA1000851	F-HEMBA1000851	0. 63
HEMBA1000869		HEMBA1000867	F-HEMBA1000867	1
HEMBA1000872 F-HEMBA1000872 0.54 HEMBA1000910 F-HEMBA1000910 0.91 HEMBA1000918 F-HEMBA1000918 0.33 HEMBA1000942 F-HEMBA1000919 0.33 HEMBA1000945 F-HEMBA1000942 0.31 HEMBA1000966 F-HEMBA1000968 0.39 HEMBA1000975 F-HEMBA1000971 0.74 HEMBA1000975 F-HEMBA1000975 0.46 HEMBA1000976 F-HEMBA1000975 0.46 HEMBA1001008 F-HEMBA1000976 0.86 HEMBA1001008 F-HEMBA1001008 0.47 HEMBA1001008 F-HEMBA1001008 0.47 HEMBA100102 F-HEMBA1001022 0.57 HEMBA1001043 F-HEMBA1001052 0.83 HEMBA1001055 F-HEMBA1001052 0.83 HEMBA1001056 F-HEMBA1001059 0.78 HEMBA1001080 F-HEMBA1001080 0.77 HEMBA1001080 F-HEMBA1001080 0.77 HEMBA1001108 F-HEMBA1001080 0.77 HEMBA1001109 F-HEMBA1001080 0.77 HEMBA1001109 F-HEMBA1001080 0.77 HEMBA1001107 F-HEMBA1001080 0.77 HEMBA1001107 F-HEMBA1001080 0.90 HEMBA1001107 F-HEMBA1001103 0.45 HEMBA1001107 F-HEMBA1001103 0.45 HEMBA1001107 F-HEMBA1001103 0.45 HEMBA1001107 F-HEMBA1001103 0.66 HEMBA1001107 F-HEMBA1001103 0.61 HEMBA1001127 F-HEMBA1001123 0.61 HEMBA1001287 F-HEMBA1001235 0.94 HEMBA1001287 F-HEMBA1001289 0.94 HEMBA1001287 F-HEMBA1001280 0.94 HEMBA1001301 F-HEMBA1001303 0.79 HEMBA1001301 F-HEMBA1001303 0.79 HEMBA1001301 F-HEMBA1001303 0.65 HEMBA1001301 F-HEMBA1001303 0.79 HEMBA1001301 F-HEMBA1001303 0.79 HEMBA1001301 F-HEMBA1001303 0.65 HEMBA1001301 F-HEMBA1001303 0.65 HEMBA1001301 F-HEMBA1001303 0.69 HEMBA1001301 F-HEMBA1001301 0.66	5	HEMBA1000869	F-HEMBA1000869	0. 82
HEMBA1000910		· · •		
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HEMBA1001133 F-HEMBA1001133 0.45 HEMBA1001137 F-HEMBA1001137 0.37 HEMBA1001140 F-HEMBA1001140 0.43 HEMBA1001174 F-HEMBA1001174 0.44 HEMBA1001197 F-HEMBA1001197 0.65 HEMBA1001213 F-HEMBA1001213 0.61 HEMBA1001235 F-HEMBA1001235 0.94 HEMBA1001247 F-HEMBA1001247 0.31 HEMBA1001257 F-HEMBA1001257 0.92 HEMBA1001281 F-HEMBA1001281 0.94 HEMBA1001286 F-HEMBA1001280 0.49 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001305 F-HEMBA1001310 0.66 HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001310 F-HEMBA1001311 0.65 HEMBA1001311 F-HEMBA1001351 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001377 F-HEMBA1001377 0.87	30	HEMBA1001109	F-HEMBA1001109	0.50
HEMBA1001137 F-HEMBA1001137 0.37 HEMBA1001140 F-HEMBA1001140 0.43 HEMBA1001174 F-HEMBA1001174 0.44 HEMBA1001197 F-HEMBA1001197 0.65 HEMBA1001213 F-HEMBA1001213 0.61 HEMBA1001235 F-HEMBA1001235 0.94 HEMBA1001247 F-HEMBA1001247 0.31 HEMBA1001257 F-HEMBA1001257 0.92 HEMBA1001281 F-HEMBA1001281 0.94 HEMBA1001286 F-HEMBA1001286 0.49 HEMBA1001289 F-HEMBA1001289 0.94 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001351 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001377 F-HEMBA1001377 0.94		HEMBA1001122	F-HEMBA1001122	0. 74
HEMBA1001140 F-HEMBA1001140 0.43 HEMBA1001174 F-HEMBA1001174 0.44 HEMBA1001197 F-HEMBA1001197 0.65 HEMBA1001213 F-HEMBA1001213 0.61 HEMBA1001235 F-HEMBA1001235 0.94 HEMBA1001247 F-HEMBA1001247 0.31 HEMBA1001257 F-HEMBA1001257 0.92 HEMBA1001281 F-HEMBA1001281 0.94 HEMBA1001286 F-HEMBA1001280 0.49 HEMBA1001289 F-HEMBA1001289 0.94 HEMBA1001302 F-HEMBA1001302 0.62 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001305 F-HEMBA1001310 0.66 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001377 F-HEMBA1001377 0.94		HEMBA1001133	F-HEMBA1001133	0.45
HEMBA1001174 F-HEMBA1001174 0. 44 HEMBA1001197 F-HEMBA1001197 0. 65 HEMBA1001213 F-HEMBA1001213 0. 61 HEMBA1001235 F-HEMBA1001235 0. 94 HEMBA1001247 F-HEMBA1001247 0. 31 HEMBA1001257 F-HEMBA1001257 0. 92 HEMBA1001281 F-HEMBA1001281 0. 94 HEMBA1001286 F-HEMBA1001280 0. 49 HEMBA1001302 F-HEMBA1001303 0. 79 HEMBA1001303 F-HEMBA1001303 0. 79 HEMBA1001304 F-HEMBA1001310 0. 66 HEMBA1001310 F-HEMBA1001310 0. 65 HEMBA1001351 F-HEMBA1001351 0. 65 HEMBA1001361 F-HEMBA1001361 0. 94 HEMBA1001377 F-HEMBA1001377 0. 87 HEMBA1001377 F-HEMBA1001377 0. 94		HEMBA1001137	F-HEMBA1001137	0. 37
HEMBA1001174 F-HEMBA1001174 0. 44 HEMBA1001197 F-HEMBA1001197 0. 65 HEMBA1001213 F-HEMBA1001213 0. 61 HEMBA1001235 F-HEMBA1001235 0. 94 HEMBA1001247 F-HEMBA1001247 0. 31 HEMBA1001257 F-HEMBA1001257 0. 92 HEMBA1001281 F-HEMBA1001281 0. 94 HEMBA1001286 F-HEMBA1001280 0. 49 HEMBA1001302 F-HEMBA1001289 0. 94 HEMBA1001303 F-HEMBA1001303 0. 79 HEMBA1001303 F-HEMBA1001310 0. 66 HEMBA1001310 F-HEMBA1001310 0. 65 HEMBA1001351 F-HEMBA1001351 0. 65 HEMBA1001377 F-HEMBA1001377 0. 87 HEMBA1001377 F-HEMBA1001377 0. 94	35	HEMBA1001140	F-HEMBA1001140	0. 43
HEMBA1001213 F-HEMBA1001213 0.61 HEMBA1001235 F-HEMBA1001235 0.94 HEMBA1001247 F-HEMBA1001247 0.31 HEMBA1001257 F-HEMBA1001257 0.92 HEMBA1001281 F-HEMBA1001281 0.94 HEMBA1001286 F-HEMBA1001286 0.49 HEMBA1001302 F-HEMBA1001303 0.79 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94		HEMBA1001174	F-HEMBA1001174	0. 44
HEMBA1001235 F-HEMBA1001235 0.94 HEMBA1001247 F-HEMBA1001247 0.31 HEMBA1001257 F-HEMBA1001257 0.92 HEMBA1001281 F-HEMBA1001281 0.94 HEMBA1001286 F-HEMBA1001286 0.49 HEMBA1001302 F-HEMBA1001302 0.62 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94		HEMBA1001197	F-HEMBA1001197	0. 65
HEMBA1001247 F-HEMBA1001247 0.31 HEMBA1001257 F-HEMBA1001257 0.92 HEMBA1001281 F-HEMBA1001281 0.94 HEMBA1001286 F-HEMBA1001286 0.49 HEMBA1001302 F-HEMBA1001302 0.62 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94		HEMBA1001213	F-HEMBA1001213	0. 61
HEMBA1001247 F-HEMBA1001247 0.31 HEMBA1001257 F-HEMBA1001257 0.92 HEMBA1001281 F-HEMBA1001281 0.94 HEMBA1001286 F-HEMBA1001286 0.49 HEMBA1001302 F-HEMBA1001302 0.62 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 50 HEMBA1001326 F-HEMBA1001326 0.59 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001377 0.87 HEMBA1001377 F-HEMBA1001377 0.87	40	HEMBA1001235	F-HEMBA1001235	0.94
HEMBA1001281 F-HEMBA1001281 0.94 HEMBA1001286 F-HEMBA1001286 0.49 HEMBA1001289 F-HEMBA1001289 0.94 HEMBA1001302 F-HEMBA1001302 0.62 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94	40	HEMBA1001247	F-HEMBA1001247	0.31
HEMBA1001286 F-HEMBA1001286 0.49 HEMBA1001289 F-HEMBA1001289 0.94 HEMBA1001302 F-HEMBA1001302 0.62 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 MEMBA1001326 F-HEMBA1001326 0.59 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94		HEMBA1001257	F-HEMBA1001257	0. 92
HEMBA1001289 F-HEMBA1001289 0.94 HEMBA1001302 F-HEMBA1001302 0.62 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 MEMBA1001326 F-HEMBA1001326 0.59 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94		HEMBA1001281	F-HEMBA1001281	0.94
HEMBA1001302 F-HEMBA1001302 0.62 HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001326 F-HEMBA1001326 0.59 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94		HEMBA1001286	F-HEMBA1001286	0.49
HEMBA1001303 F-HEMBA1001303 0.79 HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001326 F-HEMBA1001326 0.59 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94	45	HEMBA1001289	F-HEMBA1001289	0.94
HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001326 F-HEMBA1001326 0.59 HEMBA1001351 F-HEMBA1001361 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94		HEMBA1001302	F-HEMBA1001302	0. 62
HEMBA1001310 F-HEMBA1001310 0.66 HEMBA1001326 F-HEMBA1001326 0.59 HEMBA1001351 F-HEMBA1001361 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94		HEMBA1001303	F-HEMBA1001303	0. 79
HEMBA1001326 F-HEMBA1001326 0.59 HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94				
HEMBA1001351 F-HEMBA1001351 0.65 HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94	50	HEMBA1001326	F-HEMBA1001326	i
HEMBA1001361 F-HEMBA1001361 0.94 HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94		HEMBA1001351	F-HEMBA1001351	1
HEMBA1001377 F-HEMBA1001377 0.87 HEMBA1001387 F-HEMBA1001387 0.94				
55 HEMBA1001387 F-HEMBA1001387 0.94				
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	HEMBA1001398	F-HEMBA1001398	0.74
	HEMBA1001405	F-HEMBA1001405	0.61
	HEMBA1001407	F-HEMBA1001407	0. 43
5	HEMBA1001413	F-HEMBA1001413	0.40
	HEMBA1001415	F-HEMBA1001415	0.94
	HEMBA1001446	F-HEMBA1001446	0.90
	HEMBA1001450	F-HEMBA1001450	0.40
10	HEMBA1001455	F-HEMBA1001455	0. 73
	HEMBA1001476	F-HEMBA1001476	0. 65
	HEMBA1001497	F-HEMBA1001497	0. 48
	HEMBA1001510	F-HEMBA1001510	0.36
15	HEMBA1001526	F-HEMBA1001526	0.66
	HEMBA1001533	F-HEMBA1001533	0. 46
	HEMBA1001569	F-HEMBA1001569,	0.73
	HEMBA1001570	F-HEMBA1001570	0.53
20 .	HEMBA1001579	F-HEMBA1001579	0.82
	HEMBA1001581	F-HEMBA1001581	0.80
	HEMBA1001595	F-HEMBA1001595	0.72
	HEMBA1001620	F-HEMBA1001620	0.94
25	HEMBA1001635	F-HEMBA1001635	0.87
	HEMBA1001640	F-HEMBA1001640	0.82
	HEMBA1001647	F-HEMBA1001647	0.91
	HEMBA1001655	F-HEMBA1001655	0.39
30	HEMBA1001661	F~HEMBA1001661	0.86
	HEMBA1001672	F-HEMBA1001672	0.94
	HEMBA1001702	F-HEMBA1001702	0.72
	HEMBA1001711	F-HEMBA1001711	0.74
35	HEMBA1001714	F-HEMBA1001714	0.89
-	HEMBA1001723	F-HEMBA1001723	0.94
	HEMBA1001731	F-HEMBA1001731	0.37
	HEMBA1001744	F-HEMBA1001744	0.31
40	HEMBA1001746	F-HEMBA1001746	0.39
,,	HEMBA1001781	F-HEMBA1001781	0.67
	HEMBA1001800	F-HEMBA1001800	0.69
	HEMBA1001804	F-HEMBA1001804	0. 85
45	HEMBA1001809	F-HEMBA1001809	0. 78
40	HEMBA1001815	F-HEMBA1001815	0.49
	HEMBA1001819	F-HEMBA1001819	0.90
	HEMBA1001822	F-HEMBA1001822	0.80
50	HEMBA1001824	F-HEMBA1001824	0.68
30	HEMBA1001847	F-HEMBA1001847	0.46
	HEMBA1001864	F-HEMBA1001864	0.34
	HEMBA1001866	F-HEMBA1001866	0, 53
55	HEMBA1001869	F-HEMBA1001869	0. 69
55	HEMBA1001896	F-HEMBA1001896	0.48
	HEMBA1001910	F-HEMBA1001910	0. 84

			1
	HEMBA1001912	F-HEMBA1001912	0.48
	HEMBA1001913	F-HEMBA1001913	0.94
	HEMBA1001915	F-HEMBA1001915	0.41
5	HEMBA1001921	F-HEMBA1001921	0.94
	HEMBA1001939	F-HEMBA1001939	0.87
	HEMBA1001950	F-HEMBA1001950	0.36
	HEMBA1001967	F-HEMBA1001967	0.94
10	HEMBA1001987	F-HEMBA1001987	0.41
	HEMBA1002018	F-HEMBA1002018	0.41
	HEMBA1002035	F-HEMBA1002035	0.36
	HEMBA1002049	F-HEMBA1002049	0. 39
15	HEMBA1002084	F-HEMBA1002084	0.45
7.0	HEMBA1002004	F-HEMBA1002092	0. 94
	HEMBA1002092	F-HEMBA1002092	
	HEMBA1002102	F-HEMBA1002102,	0. 60
		F-HEMBA1002119	0. 88
20	HEMBA1002125		0. 31
	HEMBA1002150	F-HEMBA1002150	0. 31
	HEMBA1002151	F-HEMBA1002151	0. 82
	HEMBA1002161	F-HEMBA1002161	
25	HEMBA1002177	F-HEMBA1002177	0.38
	HEMBA1002189	F-HEMBA1002189	0.43
	HEMBA1002191	F-HEMBA1002191	0.94
	HEMBA1002199	F-HEMBA1002199	0.85
30	HEMBA1002212	F-HEMBA1002212	0.39
	HEMBA1002215	F-HEMBA1002215	0. 94
	HEMBA1002229	F-HEMBA1002229	0.83
	HEMBA1002237	F-HEMBA1002237	0.86
35	HEMBA1002241	F-HEMBA1002241	0.66
	HEMBA1002265	F-HEMBA1002265	0.38
	HEMBA1002267	F-HEMBA1002267	0. 33
	HEMBA1002341	F-HEMBA1002341	0.53
40	HEMBA1002363	F-HEMBA1002363	0.88
	HEMBA1002417	F-HEMBA1002417	0. 83
	HEMBA1002419	F-HEMBA1002419	0. 92
	HEMBA1002430	F-HEMBA1002430	0. 45
45	HEMBA1002439	F-HEMBA1002439	0. 83
45	HEMBA1002458	F-HEMBA1002458	0.47
	HEMBA1002460	F-HEMBA1002460	0.48
	HEMBA1002462	F-HEMBA1002462	0. 55
	HEMBA1002469		1
50	HEMBA1002475	F-HEMBA1002475	0. 88
	HEMBA1002477	F-HEMBA1002477	0. 45
	HEMBA1002495		0. 78
	HEMBA1002503	F-HEMBA1002503	0. 94
55	HEMBA1002508		0. 33
	HEMBA1002513	F-HEMBA1002513	0. 41

HEMBA1002542 F-HEMBA1002542 0.4 HEMBA1002547 F-HEMBA1002547 0.4 HEMBA1002569 F-HEMBA1002569 0.8	5
5 HEMBA1002569 F-HEMBA1002569 0.8	5
LICHION TOOS OF A LICHION TOOS OF O	
HEMBA1002583 F-HEMBA1002583 0.8	,
HEMBA1002609 F-HEMBA1002609 0.7	5
HEMBA1002624 F-HEMBA1002624 0.5	9
10 HEMBA1002688 F-HEMBA1002688 0.8	3
HEMBA1002696 F-HEMBA1002696 0.8	2
HEMBA1002703 F-HEMBA1002703 0.9	4
HEMBA1002746 F-HEMBA1002746 0.3	3
15 HEMBA1002750 F-HEMBA1002750 0.5	3
HEMBA1002768 F-HEMBA1002768 0.9	4
HEMBA1002770 F-HEMBA1002770, 0.6	1
HEMBA1002777 F-HEMBA1002777 0.9	0
20 HEMBA1002779 F-HEMBA1002779 0.9	2
HEMBA1002794 F-HEMBA1002794 0.8	6
HEMBA1002810 F-HEMBA1002810 0.8	7
HEMBA1002816 F-HEMBA1002816 0.9	1
25 HEMBA1002818 F-HEMBA1002818 0.9	4
HEMBA1002850 F-HEMBA1002850 0.5	7
HEMBA1002863 F-HEMBA1002863 0.4	В
HEMBA1002876 F-HEMBA1002876 0.8	6
30 HEMBA1002935 F-HEMBA1002935 0.9	1
HEMBA1002937 F-HEMBA1002937 0.6	1
HEMBA1002939 F-HEMBA1002939 0.3	2
HEMBA1002951 F-HEMBA1002951 0.8	3
35 HEMBA1002954 F-HEMBA1002954 0. 6	
HEMBA1002970 F-HEMBA1002970 0.3	
HEMBA1002971 F-HEMBA1002971 0.5	
HEMBA1002973 F-HEMBA1002973 0.3	
40 HEMBA1002997 F-HEMBA1002997 0.4	
HEMBA1002999 F-HEMBA1002999 0.4	
HEMBA1003021 F-HEMBA1003021 0.7	
HEMBA1003033 F-HEMBA1003033 0.9	
HEMBA1003035 F-HEMBA1003035 0.6	
HEMBA1003041 F-HEMBA1003041 0.4	-
HEMBA1003046 F-HEMBA1003046 0.9	
HEMBA1003067 F-HEMBA1003067 0.3	
HEMBA1003077 F-HEMBA1003077 0.8	
NEMBATUUSU/8 F-NEMBATUUSU/8 U. 8	
HEMBA1003079 F-HEMBA1003079 0.4	
HEMBA1003096 F-HEMBA1003096 0.3	
HEMBA1003117 F-HEMBA1003117 0.9 55 HEMBA1003129 F-HEMBA1003129 0.8	
11cmDA1003123 1 11cmDA1000123 0. 0	
HEMBA1003136 F-HEMBA1003136 0.7	'

	HEMBA1003148	F-HEMBA1003148	0. 91
	HEMBA1003175	F-HEMBA1003175	0. 41
	HEMBA1003179	F-HEMBA1003179	0. 42
5	HEMBA1003199	F-HEMBA1003199	0, 84
	HEMBA1003222	F-HEMBA1003222	0. 47
	HEMBA1003235	F-HEMBA1003235	0.94
	HEMBA1003250	F-HEMBA1003250	0. 39
10	HEMBA1003257	F-HEMBA1003257	0.88
•	HEMBA1003281	F-HEMBA1003281	0. 67
	HEMBA1003286	F-HEMBA1003286	0. 57
	HEMBA1003291	F-HEMBA1003291	0.89
15	HEMBA1003322	F-HEMBA1003322	0. 60
	HEMBA1003327	F-HEMBA1003327	0.88
	HEMBA1003369	F-HEMBA1003369,	
	HEMBA1003370	F-HEMBA1003370	0.75
20	HEMBA1003380	F-HEMBA1003380	0.68
	HEMBA1003395	F-HEMBA1003395	0.88
	HEMBA1003402	F-HEMBA1003402	0. 45
	HEMBA1003403	F-HEMBA1003403	0. 44
25	HEMBA1003408	F-HEMBA1003408	0.34
	HEMBA1003417	F-HEMBA1003417	0. 78
	HEMBA1003418	F-HEMBA1003418	0, 94
	HEMBA1003433	F-HEMBA1003433	0. 70
30	HEMBA1003447	F-HEMBA1003447	0. 49
	HEMBA1003461	F-HEMBA1003461	0. 85
	HEMBA1003463	F-HEMBA1003463	0. 59
	HEMBA1003480	F-HEMBA1003480	0. 94
35	HEMBA1003528	F-HEMBA1003528	0. 90
	HEMBA1003538	F-HEMBA1003538	0. 47
	HEMBA1003545	F-HEMBA1003545	0. 74
	HEMBA1003555	F-HEMBA1003555	0.94
40	HEMBA1003556	F-HEMBA1003556	0. 44
	HEMBA1003560	F-HEMBA1003560	0. 40
	HEMBA1003568	F-HEMBA1003568	0. 94
	HEMBA1003569	F-HEMBA1003569	0. 82
45	HEMBA1003581	F-HEMBA1003581	0. 83
	HEMBA1003591	F-HEMBA1003591	0. 68
	HEMBA1003615	F-HEMBA1003615	0.94
	HEMBA1003617	F-HEMBA1003617	0.88
50	HEMBA1003621 HEMBA1003645	F-HEMBA1003621 F-HEMBA1003645	0.47
	HEMBA1003646	F-HEMBA1003646	0. 75 0. 51
	HEMBA1003662	F-HEMBA1003662	0. 51
	HEMBA1003667	F-HEMBA1003667	0. 54
55	HEMBA1003679	F-HEMBA1003679	0. 72
	HEMBA1003680	F-HEMBA1003680	0. 72
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	HEMBA1003684	F-HEMBA1003684	0. 94
	HEMBA1003690	F-HEMBA1003690	0.93
	HEMBA1003692	F-HEMBA1003692	0.46
5	HEMBA1003711	F-HEMBA1003711	0. 94
	HEMBA1003720	F-HEMBA1003720	0. 94
	HEMBA1003729	F-HEMBA1003729	0.73
	HEMBA1003742	F-HEMBA1003742	0. 94
10	HEMBA1003760	F-HEMBA1003760	0.75
	HEMBA1003783	F-HEMBA1003783	0.31
	HEMBA1003799	F-HEMBA1003799	0, 70
	HEMBA1003803	F-HEMBA1003803	0.87
15	HEMBA1003804	F-HEMBA1003804	0.94
	HEMBA1003805	F-HEMBA1003805	0.94
	HEMBA1003807	F-HEMBA1003807,	
	HEMBA1003827	F-HEMBA1003827	0. 62
20	HEMBA1003836	F-HEMBA1003836	0.69
	HEMBA1003838	F-HEMBA1003838	0.80
	HEMBA1003864	F-HEMBA1003864	0.94
	HEMBA1003866	F-HEMBA1003866	0.32
25	HEMBA1003879	F-HEMBA1003879	0.82
20	HEMBA1003880	F-HEMBA1003880	0. 59
	HEMBA1003893	F-HEMBA1003893	0.47
	HEMBA1003953	F-HEMBA1003953	0. 77
30	HEMBA1003959	F-HEMBA1003959	0.32
30	HEMBA1003978	F-HEMBA1003978	0. 37
	HEMBA1003985	F-HEMBA1003985	0. 31
	HEMBA1003989	F-HEMBA1003989	0. 75
35	HEMBA1004011	F-HEMBA1004011	0. 34
35	HEMBA1004048	F-HEMBA1004048	0. 32
	HEMBA1004055	F-HEMBA1004055	0.74
	HEMBA1004056	F-HEMBA1004056	0. 49
40	HEMBA1004074	F-HEMBA1004074	0. 39
40	HEMBA1004086	F-HEMBA1004086	0. 45
	HEMBA1004097	F-HEMBA1004097	0. 82
	HEMBA1004111	F-HEMBA1004111	0. 94
4-	HEMBA1004131	F-HEMBA1004131	0. 84
45	HEMBA1004133	F-HEMBA1004133	0. 40
	HEMBA1004143	F-HEMBA1004143	0. 33
	HEMBA1004146	F-HEMBA1004146	0. 31
	HEMBA1004150	F-HEMBA1004150	0. 70
50	HEMBA1004168	F-HEMBA1004168	0. 34
	HEMBA1004199	F-HEMBA1004199	0. 34
•	HEMBA1004200	F-HEMBA1004200	0, 40
	HEMBA1004202	F-HEMBA1004202	0. 88
55	HEMBA1004203	F-HEMBA1004203	0. 94
	HEMBA1004207	F-HEMBA1004207	0. 53

			1
	HEMBA1004227	F-HEMBA1004227	0.94
	HEMBA1004238	F-HEMBA1004238	0. 44
	HEMBA1004246	F-HEMBA1004246	0.71
5	HEMBA1004248	F-HEMBA1004248	0.40
	HEMBA1004264	F-HEMBA1004264	0. 94
	HEMBA1004274	F-HEMBA1004274	0. 88
	HEMBA1004275	F-HEMBA1004275	0. 85
10	HEMBA1004276	F-HEMBA1004276	0. 32
	HEMBA1004286	F-HEMBA1004286	0. 37
	HEMBA1004289	F-HEMBA1004289	0. 76
	HEMBA1004321	F-HEMBA1004321	0. 82
15	HEMBA1004327	F-HEMBA1004327	0. 40
	HEMBA1004330	F-HEMBA1004330	0. 40
	HEMBA1004335	F-HEMBA1004335,	- 1
	HEMBA1004341	F-HEMBA1004331	0. 73
20	HEMBA1004341		0. 79
	HEMBA1004372		0. 79
	HEMBA1004372		0. 32
	HEMBA1004389		0. 70
25	HEMBA1004479		0. 75
	HEMBA1004479	F-HEMBA1004479	0. 62
	HEMBA1004502	F-HEMBA1004502	0. 02
	HEMBA1004507	F-HEMBA1004507	0. 71
30	HEMBA1004509	F-HEMBA1004509	0. 68
30	HEMBA1004534		0. 56
	HEMBA1004542		0.36
	HEMBA1004554		0. 90
25	HEMBA1004560		0.57
35	HEMBA1004573		0. 93
	HEMBA1004596	F-HEMBA1004596	0. 45
	HEMBA1004604		0. 91
	HEMBA1004610	F-HEMBA1004610	0. 59
40	HEMBA1004632	F-HEMBA1004632	0. 59
	HEMBA1004637	F-HEMBA1004637	0. 38
	HEMBA1004638	F-HEMBA1004638	0. 40
	HEMBA1004669	F-HEMBA1004669	0. 94
45	HEMBA1004693	F-HEMBA1004693	0. 52
	HEMBA1004697	F-HEMBA1004697	0. 94
	HEMBA1004705	F-HEMBA1004705	0.94
	HEMBA1004709	F-HEMBA1004709	0. 77
50	HEMBA1004703	F-HEMBA1004701	0. 77
	HEMBA1004711	F-HEMBA1004711	0. 73
	HEMBA1004734	F-HEMBA1004734	0. 94
	HEMBA1004736	F-HEMBA1004736	0. 94
55	HEMBA1004751	F-HEMBA1004751	0. 64
	HEMBA1004752	F-HEMBA1004752	0. 87
_	HUNDA1004/32	1 11LRDA1004732	0. 67

	HEMBA1004753	F-HEMBA1004753 ·	0. 54
	HEMBA1004756	F-HEMBA1004756	0. 54
	HEMBA1004758	F-HEMBA1004758	- 1
5	HEMBA1004763	F-HEMBA1004763	
	HEMBA1004768	F-HEMBA1004768	0.66
	HEMBA1004771	F-HEMBA1004771	0. 72
	HEMBA1004776	F-HEMBA1004776	0. 49
10	HEMBA1004795	F-HEMBA1004795	0. 46
	HEMBA1004806	F-HEMBA1004806	0. 32
	HEMBA1004847	F-HEMBA1004847	0. 79
	HEMBA1004850	F-HEMBA1004850	0. 94
15	HEMBA1004863	F-HEMBA1004863	0. 32
	HEMBA1004864	F-HEMBA1004864	0.94
	HEMBA1004889	F-HEMBA1004889,	
	HEMBA1004923	F-HEMBA1004923	0.70
20	HEMBA1004929	F-HEMBA1004929	0.31
	HEMBA1004930	F-HEMBA1004930	0.90
	HEMBA1004933	F-HEMBA1004933	0.84
•	HEMBA1004934	F-HEMBA1004934	0.34
25	HEMBA1004944	F-HEMBA1004944	0.40
	HEMBA1004954	F~HEMBA1004954	0. 62
	HEMBA1004972	F-HEMBA1004972	0. 90
	HEMBA1004973	F~HEMBA1004973	0. 94
30	HEMBA1004977	F-HEMBA1004977	0.77
	HEMBA1004980	F-HEMBA1004980	0. 33
	HEMBA1005009	F-HEMBA1005009	0.94
	HEMBA1005019	F-HEMBA1005019	0. 32
35	HEMBA1005029	F-HEMBA1005029	0.84
	HEMBA1005035	F-HEMBA1005035	0.91
	HEMBA1005047	F-HEMBA1005047	0.94
	HEMBA1005050	F-HEMBA1005050	0. 88
40	HEMBA1005066	F-HEMBA1005066	0. 79
	HEMBA1005075	F-HEMBA1005075	0.31
	HEMBA1005079	F-HEMBA1005079	0. 35
	HEMBA1005083	F-HEMBA1005083	0. 43
45	HEMBA1005101	F-HEMBA1005101	0. 94
40	HEMBA1005113	F-HEMBA1005113	0. 38
	HEMBA1005133	F-HEMBA1005133	0. 34
	HEMBA1005149		0. 36
50	HEMBA1005185	F-HEMBA1005185	0. 64
30	HEMBA1005201	F-HEMBA1005201	0. 83
	HEMBA1005202	F-HEMBA1005202	0. 94
	HEMBA1005206	F-HEMBA1005206	0. 94
EE	HEMBA1005219	F-HEMBA1005219	0. 57
55	HEMBA1005223	F-HEMBA1005223	0. 33
	HEMBA1005244	F-HEMBA1005244	0. 35
			

	HEMBA1005252	F-HEMBA1005252	0. 80
	HEMBA1005296	F-HEMBA1005296	0. 94
	HEMBA1005314	F-HEMBA1005314	0. 94
5	HEMBA1005331	F-HEMBA1005331	0. 69
	HEMBA1005338	F-HEMBA1005338	0. 94
	HEMBA1005367	F-HEMBA1005367	0. 50
	HEMBA1005382	F-HEMBA1005382	0. 91
10	HEMBA1005394	F-HEMBA1005394	0. 45
	HEMBA1005403	F-HEMBA1005403	0. 94
	HEMBA1005411	F-HEMBA1005411	0. 33
	HEMBA1005423	F-HEMBA1005423	0. 64
15	HEMBA1005468	F-HEMBA1005468	0. 33
	HEMBA1005469	F-HEMBA1005469	0. 59
	HEMBA1005472	F-HEMBA1005472,	
	HEMBA1005474	F-HEMBA1005474	0. 51
20	HEMBA1005475	F-HEMBA1005475	0. 36
	HEMBA1005513	F-HEMBA1005513	0. 94
	HEMBA1005517	F-HEMBA1005517	0. 94
	HEMBA1005518	F-HEMBA1005518	0. 39
25	HEMBA1005526	F-HEMBA1005526	0. 62
	HEMBA1005528	F-HEMBA1005528	0.81
	HEMBA1005530	F-HEMBA1005530	0. 62
	HEMBA1005548	F-HEMBA1005548	0.94
30	HEMBA1005558	F-HEMBA1005558	0.94
	HEMBA1005576	F-HEMBA1005576	0. 49
	HEMBA1005581	F-HEMBA1005581	0. 53
	HEMBA1005582	F-HEMBA1005582	0. 34
35	HEMBA1005583	F-HEMBA1005583	0. 94
-	HEMBA1005595	F-HEMBA1005595	0. 93
	HEMBA1005609	F-HEMBA1005609	0. 34
	HEMBA1005621	F-HEMBA1005621	0. 94
40	HEMBA1005666	F-HEMBA1005666	0. 57
	HEMBA1005680	F-HEMBA1005680	0. 38
	HEMBA1005685	F-HEMBA1005685	0. 32
	HEMBA1005732	F-HEMBA1005732	0. 33
45	HEMBA1005737	F-HEMBA1005737	0. 92
10	HEMBA1005746	F-HEMBA1005746	0. 37
	HEMBA1005755	F-HEMBA1005755	0.94
	HEMBA1005813	F-HEMBA1005813	0. 33
50	HEMBA1005815	F-HEMBA1005815	0.82
	HEMBA1005822	F-HEMBA1005822	0.83
	HEMBA1005834 HEMBA1005852	F-HEMBA1005834 F-HEMBA1005852	0. 70 0. 76
	HEMBA1005884		0. 7 0 0. 79
55	HEMBA1005891	F-HEMBA1005884 F-HEMBA1005891	0. 79 0. 94
	HEMBA1005963	F-HEMBA1005963	0. 40
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	HEMBA1005990	F-HEMBA1005990	0. 82
	HEMBA1005991	F-HEMBA1005991	0. 43
	HEMBA1006031	F-HEMBA1006031	0. 55
5	HEMBA1006035	F-HEMBA1006035	0. 43
	HEMBA1006081	F-HEMBA1006081	0. 36
	HEMBA1006091	F-HEMBA1006091	0. 81
	HEMBA1006100	F-HEMBA1006100	0. 31
10	HEMBA1006108	F-HEMBA1006108	0. 80
•	HEMBA1006121	F-HEMBA1006121	0. 45
	HEMBA1006130	F-HEMBA1006130	0.46
	HEMBA1006138	F-HEMBA1006138	0.48
15	HEMBA1006155	F-HEMBA1006155	0. 72
	HEMBA1006173	F-HEMBA1006173	0. 42
	HEMBA1006182	F-HEMBA1006182,	0. 73
	HEMBA1006198	F-HEMBA1006198	0. 94
20	HEMBA1006235	F-HEMBA1006235	0.58
	HEMBA1006248	F-HEMBA1006248	0. 57
	HEMBA1006252	F-HEMBA1006252	0. 31
	HEMBA1006272	F-HEMBA1006272	0. 54
25	HEMBA1006278	F-HEMBA1006278	0. 92
	HEMBA1006283	F-HEMBA1006283	0.90
	HEMBA1006284	F-HEMBA1006284	0. 93
	HEMBA1006291	F-HEMBA1006291	0.38
30	HEMBA1006293	F-HEMBA1006293	0.89
30	HEMBA1006309	F-HEMBA1006309	0. 94
	HEMBA1006310	F-HEMBA1006310	0. 67
	HEMBA1006344	F-HEMBA1006344	0.94
35	HEMBA1006347	F-HEMBA1006347	0.94
33	HEMBA1006349	F-HEMBA1006349	0.56
	HEMBA1006377	F-HEMBA1006377	0.94
	HEMBA1006381	F-HEMBA1006381	0. 33
40	HEMBA1006398	F-HEMBA1006398	0. 71
40	HEMBA1006424	F-HEMBA1006424	0. 40
	HEMBA1006445	F-HEMBA1006445	0. 86
	HEMBA1006467	F-HEMBA1006467	0. 55
ar.	HEMBA1006474	F-HEMBA1006474	0. 66
45	HEMBA1006483	F-HEMBA1006483	0. 71
	HEMBA1006485	F-HEMBA1006485	0. 87
	HEMBA1006492	F-HEMBA1006492	0. 73
	HEMBA1006494	F-HEMBA1006494	0. 90
50	HEMBA1006497	F-HEMBA1006497	0. 77
	HEMBA1006502	F-HEMBA1006502	0. 80
	HEMBA1006507	F-HEMBA1006507	0. 56
	HEMBA1006521	F-HEMBA1006521	0. 91
55	HEMBA1006530	F-HEMBA1006530	0. 37
	HEMBA1006535	F-HEMBA1006535	056

		014 011 AZ	
	HEMBA1006559	F-HEMBA1006559	0.39
	HEMBA1006566	F-HEMBA1006566	0.41
	HEMBA1006579	F-HEMBA1006579	0.83
5	HEMBA1006583	F-HEMBA1006583	0. 31
	HEMBA1006612	F-HEMBA1006612	0. 55
	HEMBA1006624	F-HEMBA1006624	0. 94
	HEMBA1006631	F-HEMBA1006631	0.74
10	HEMBA1006643	F-HEMBA1006643	0.38
	HEMBA1006652	F-HEMBA1006652	0. 93
	HEMBA1006659	F-HEMBA1006659	0. 63
	HEMBA1006674	F-HEMBA1006674	0. 54
15	HEMBA1006682	F-HEMBA1006682	0. 34
	HEMBA1006708	F-HEMBA1006708	0.40
	HEMBA1006709	F-HEMBA1006709,	
	HEMBA1006717	F-HEMBA1006717	0. 43
20	HEMBA1006737	F-HEMBA1006737	0. 72
20	HEMBA1006754	F-HEMBA1006754	0.45
	HEMBA1006758	F-HEMBA1006758	0.60
	HEMBA1006767	F-HEMBA1006767	0.61
25	HEMBA1006789	F-HEMBA1006789	0.34
20	HEMBA1006795	F-HEMBA1006795	0. 48
	HEMBA1006796	F-HEMBA1006796	0.41
	HEMBA1006807	F-HEMBA1006807	0.34
30	HEMBA1006832	F-HEMBA1006832	0.59
30	HEMBA1006877	F-HEMBA1006877	0.90
	HEMBA1006885	F-HEMBA1006885	0.80
	HEMBA1006900	F-HEMBA1006900	0.34
25	HEMBA1006914	F-HEMBA1006914	0.78
35	HEMBA1006926	F-HEMBA1006926	0. 67
	HEMBA1006936	F-HEMBA1006936	0. 65
	HEMBA1006941	F-HEMBA1006941	0.94
40	HEMBA1006973	F-HEMBA1006973	0. 64
40	HEMBA1006976	F-HEMBA1006976	0.40
	HEMBA1006993	F-HEMBA1006993	0.81
	HEMBA1006996	F-HEMBA1006996	0. 77
-	HEMBA1007002	F-HEMBA1007002	0. 72
45	HEMBA1007018	F-HEMBA1007018	0. 31
	HEMBA1007052	F-HEMBA1007052	0. 41
	HEMBA1007062	F-HEMBA1007062	0. 39
	HEMBA1007066	F-HEMBA1007066	0. 35
50	HEMBA1007080	F-HEMBA1007080	0.43
	HEMBA1007085	F-HEMBA1007085	0. 73
	HEMBA1007087	F-HEMBA1007087	0. 94
	HEMBA1007112	F-HEMBA1007112	0.49
55	HEMBA1007121	F-HEMBA1007121	0.70
	HEMBA1007149	F-HEMBA1007149	0. 94

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	HEMBA1007151	F-HEMBA1007151	0. 80
	HEMBA1007174	F-HEMBA1007174	0. 63
	HEMBA1007178	F-HEMBA1007178	0. 47
5	HEMBA1007194	F-HEMBA1007194	0. 94
	HEMBA1007203	F-HEMBA1007203	0. 48
	HEMBA1007224	F-HEMBA1007224	0. 74
	HEMBA1007243	F-HEMBA1007243	0. 94
10	HEMBA1007251	F-HEMBA1007251	0. 79
	HEMBA1007267	F-HEMBA1007267	0. 86
	HEMBA1007281	F-HEMBA1007281	0. 61
			0. 87
45	HEMBA1007300	F-HEMBA1007300	1
15	HEMBA1007301	F-HEMBA1007301	0. 38
	HEMBA1007319	F-HEMBA1007319	0. 47
	HEMBA1007320	F-HEMBA1007320,	
	HEMBA1007342	F-HEMBA1007342	0. 80
20	HEMBB1000008	F-HEMB81000008	0. 39
	HEMB81000018	F-HEMBB1000018	0. 79
	HEMBB1000024	F-HEMBB1000024	0. 62
	HEMBB1000025	F-HEMBB1000025	0. 66
25	HEMBB1000030	F-HEMBB1000030	0. 33
	HEMBB1000036	F-HEMBB1000036	0. 75
	HEMBB1000037	F-HEMBB1000037	0. 94
	HEMBB1000048	F-HEMBB1000048	0. 87
30	HEMBB1000083	F-HEMBB1000083	0.48
	HEMBB1000103	F-HEMBB1000103	0. 63
	HEMBB1000119	F-HEMBB1000119	0.94
	HEMBB1000136	F-HEMBB1000136	0. 92
35	HEMBB1000173	F-HEMBB1000173	0.46
35	HEMBB1000175	F-HEMBB1000175	0.48
	HEMBB1000198	F-HEMBB1000198	0.74
	HEMBB1000215	F-HEMBB1000215	0. 61
	HEMBB1000217	F-HEMBB1000217	0. 94
40	HEMBB1000226	F-HEMBB1000226	0. 94
	HEMBB1000240	F-HEMBB1000240	0.40
	HEMBB1000244	F-HEMBB1000244	0. 81
	HEMBB1000266	F-HEMBB1000266	0. 92
45	HEMBB1000338	F-HEMBB1000338	0. 33
	HEMBB1000339	F-HEMBB1000339	0. 40
	HEMBB1000391	F-HEMBB1000391	0. 43
	HEMBB1000438	F-HEMBB1000438	0. 79
50	HEMBB1000449	F-HEMBB1000449	0. 48
	HEMBB1000510	F-HEMBB1000510	0. 35
	HEMBB1000550	F-HEMBB1000550	0. 94
	HEMBB1000556	F-HEMB81000556	0. 89
55	HEMBB1000589	F-HEMBB1 000589	0. 77
-	HEMBB1000589	F-HEMBB1000591	!
	1111100 1000391	160000100031	0. 33

	HEMBB1000592	F-HEMBB1000592	0.77
	HEMBB1000593	F-HEMBB1000593	0. 68
	HEMBB1000623	F-HEMBB1000623	0.94
5	HEMBB1000630	F-HEMBB1000630	0, 59
	HEMBB1000631	F-HEMBB1000631	0.71
	HEMBB1000632	F-HEMBB1000632	0.38
	HEMBB1000671	F-HEMBB1000671	0.36
10	HEMBB1000673	F-HEMBB1000673	0. 54
70	HEMBB1000693	F-HEMBB1000693	0.68
	HEMBB1000705	F-HEMBB1000705	0. 44
	HEMB81000706	F-HEMBB1000706	0. 45
15	HEMBB1000725	F-HEMBB1000725	0.91
15	HEMBB1000763	F-HEMBB1000763	0.87
	HEMBB1000774	F-HEMBB1000774	0.80
	HEMBB1000781	F-HEMBB1000781	0.94
00	HEMBB1000789	F-HEMBB1000789	0.34
20	HEMBB1000807	F-HEMBB1000807	0.94
	HEMBB1000810	F-HEMBB1000810	0, 33
	HEMBB1000826	F-HEMBB1000826	0. 36
	HEMBB1000835	F-HEMBB1000835	0. 81
25	HEMB81000848	F-HEMBB1000848	0.66
	HEMBB1000852	F-HEMB81000852	0.85
	HEMBB1000870	F-HEMBB1000870	0.34
	HEMBB1000887	F-HEMB81000887	0.58
30	HEMBB1000908	F-HEMBB1000908	0.31
	HEMBB1000927	F-HEMBB1000927	0.94
	HEMB81000947	F-HEMBB1000947	0.33
	HEMBB1000973	F-HEMBB1000973	0.72
35	HEMBB1000975	F-HEMB81000975	0.68
	HEMB81000985	F-HEMBB1000985	0.62
	HEMBB1000991	F-HEMBB1000991	0.45
	HEMBB1001011	F-HEMBB1001011	0.58
40	HEMBB1001014	F-HEMBB1001014	0.86
	HEMBB1001020	F-HEMBB1001020	0. 87
	HEMBB1001024	F-HEMBB1001024	0.64
	HEMBB1001056	F-HEMBB1001056	0. 94
45	HEMBB1001058	F-HEMB81001058	0. 39
	HEMB81001068	F-HEMB81001068	0.57
	HEMBB1001096	F-HEMBB1001096	0. 48
	HEMBB1001105	F-HEMBB1001105	0. 45
50	HEMBB1001112	F-HEMBB1001112	0.47
	HEMBB1001117	F-HEMBB1001117	0. 83
	HEMBB1001126	F-HEMBB1001126	0. 76
	HEMBB1001137	F-HEMBB1001137	0. 53
55	HEMBB1001151	F-HEMBB1001151	0. 94
	HEMBB1001153	F-HEMBB1001153	0.77

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	HEMBB1001169	F-HEMBB1001169	0.39
	HEMBB1001175	F-HEMBB1001175	0. 91
	HEMBB1001182	F~HEMBB1001182	0. 94
5	HEMBB1001199	F-HEMBB1001199	0, 54
	HEMBB1001210	F-HEMBB1001210	0.88
	HEMBB1001234	F-HEMBB1001234	0. 67
	HEMBB1001242	F-HEMBB1001242	0. 33
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10	HEMBB1001288	F-HEMBB1001288	0. 33
	HEMBB1001289	F-HEMBB1001289	0.58
	HEMBB1001294	F-HEMBB1001294	0.86
	HEMBB1001314	F-HEMBB1001314	0.76
15	HEMBB1001331	F-HEMBB1001331	0. 44
	HEMBB1001339	F-HEMBB1001339	0.40
	HEMBB1001346	F-HEMBB1001346,	0.83
	HEMBB1001364	F-HEMBB1001364	0.69
20	HEMBB1001369	F-HEMBB1001369	0.47
	HEMBB1001384	- F-HEMBB1001384	0.80
	HEMBB1001387	F-HEMBB1001387	0. 36
	HEMBB1001394	F-HEMBB1001394	0. 94
25	HEMBB1001410	F-HEMBB1001410	0, 51
23	HEMBB1001426	F-HEMBB1001426	0. 80
	HEMBB1001429	F-HEMBB1001429	0. 91
	HEMBB1001449	F-HEMBB1001449	0. 31
	HEMB81001482	F-HEMBB1001482	0. 44
30	HEMB81001531	F-HEMBB1001531	0. 58
	HEMBB1001562		0. 65
		F-HEMBB1001562	-
	HEMBB1001564	F-HEMBB1001564	0. 46
35	HEMBB1001585	F-HEMB81001585	0. 74
	HEMBB1001603	F-HEMBB1001603	0. 81
	HEMBB1001635	F-HEMBB1001635	0, 46
	HEMBB1001653	F-HEMBB1001653	0. 33
40	HEMBB1001665	F-HEMBB1001665	0. 39
	HEMBB1001668	F-HEMBB1001668	0. 38
	HEMBB1001673	F-HEMBB1001673	0. 83
	HEMBB1001684	F-HEMBB1001684	0, 44
4-	HEMBB1001685	F-HEMBB1001685	0. 90
45	HEMBB1001695	F-HEMBB1001695	0. 92
	HEMBB1001707	F-HEMBB1001707	0. 51
	HEMBB1001736	F-HEMBB1001736	0. 94
	- HEMBB1001760	F-HEMBB1001760	0. 88
50	HEMBB1001785	F-HEMBB1001785	0. 74
	HEMBB1001812	F-HEMBB1001812	0. 33
	HEMBB1001816	F-HEMBB1001816	0. 41
	HEMBB1001831	F-HEMBB1001831	0. 93
55	HEMBB1001834	F-HEMBB1001834	0. 78
	HEMBB1001839	F-HEMBB1001839	0. 32
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	HEMBB1001869	F-HEMBB1001869	0. 48
	HEMBB1001872	F-HEMBB1001872	0. 93
	HEMBB1001874	F-HEMBB1001874	0. 31
5	HEMBB1001905	F-HEMBB1001905	0. 87
	HEMBB1001906	F-HEMBB1001906	0. 88
	HEMBB1001908	F-HEMBB1001908	0. 69
	HEMBB1001910	F-HEMBB1001910	0. 77
10	HEMBB1001915	F-HEMBB1001915	0. 76
	HEMBB1001945	F-HEMBB1001945	0. 64
	HEMBB1001947	F-HEMBB1001947	0. 94
	HEMBB1001950	F-HEMBB1001950	0. 73
15	HEMBB1001957	F-HEMBB1001957	0. 64
	HEMBB1001962	F-HEMBB1001962	0. 90
	HEMBB1001990	F-HEMBB1001990,	0. 83
	HEMBB1002044	F~HEMBB1002044	0. 82
20	HEMBB1002050	F-HEMBB1002050	0. 91
20	HEMBB1002068	F-HEMBB1002068	0. 48
	HEMBB1002134	F-HEMBB1002134	0. 80
	HEMBB1002142	F-HEMBB1002142	0. 39
25	HEMBB1002152	F-HEMBB1002152	0. 31
20	HEMBB1002193	F-HEMBB1002193	0. 35
	HEMBB1002217	F-HEMBB1002217	0.49
	HEMBB1002218	F-HEMBB1002218	0. 32
30	HEMBB1002249	F-HEMBB1002249	0. 72
30	HEMBB1002300	F-HEMBB1002300	0. 75
	HEMBB1002327	F-HEMBB1002327	0.86
	HEMBB1002329	F-HEMBB1002329	0.31
25	HEMBB1002340	F-HEMBB1002340	0. 42
35	HEMBB1002342	F-HEMBB1002342	0. 76
	HEMBB1002358	F-HEMBB1002358	0. 70
	HEMBB1002359	F-HEMBB1002359	0. 75
40	HEMBB1002383	F-HEMBB1002383	0. 78
40	HEMBB1002409	F-HEMBB1002409	0. 52
	HEMBB1002415	F-HEMBB1002415	0.82
	HEMBB1002442	F-HEMBB1002442	0. 31
	HEMBB1002457	F-HEMBB1002457	0. 46
45	HEMBB1002489	F-HEMBB1002489	0. 69
	HEMBB1002492	F-HEMBB1002492	0. 68
	HEMBB1002495	F-HEMBB1002495	0. 94
	HEMB81002502	F-HEMBB1002502	0. 39
50	HEMB81002510	F-HEMBB1002510	0. 94
	HEMBB1002545	F-HEMBB1002545	0. 44
	HEMBB1002550	F-HEMBB1002550	0.57
	HEMBB1002600	F-HEMBB1002600	0.91
55	HEMBB1002607	F-HEMBB1002607	0. 83
•	HEMBB1002614	F-HEMBB1002614	0. 75

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	HEMBB1002684	F-HEMBB1002684	0.84
	HEMBB1002692	F-HEMBB1002692	0.52
	HEMBB1002697	F-HEMBB1002697	0. 86
5	HEMBB1002705	F-HEMBB1002705	0. 32
	MAMMA1000019	F-MAMMA1000019	0. 48
	MAMMA1000020	F-MAMMA1000020	0.36
	MAMMA1000025	F-MAMMA1000025	0. 31
10	MAMMA1000055		0. 39
	MAMMA1000057		0. 35
	MAMMA1000069		0. 78
	MAMMA1000084		0. 61
15	MAMMA1000084		0. 72
15	MAMMA1000139		0. 72
	MAMMA1000133		
	MAMMA1000143		0. 40
20	MAMMA1000171	F-MAMMA1000171	0. 78
		F-MAMMA1000173	0.94
	MAMMA1000183		0.77
		F-MAMMA1000251	0.94
25	MAMMA1000257		0. 75
	MAMMA1000277		0.94
	MAMMA1000278		0. 35
	MAMMA1000279		0. 62
30	MAMMA1000284		0. 35
	MAMMA1000309		0. 53
		F-MAMMA1000312	0.68
		F-MAMMA1000313	0. 32
35		F-MAMMA1000339	0. 35
	MAMMA1000361	F-MAMMA1000361	0. 61
		F-MAMMA1000372	0. 60
	MAMMA1000388	F-MAMMA1000388	0. 93
40	MAMMA1000395	F-MAMMA1000395	0. 94
40	MAMMA1000410	F-MAMMA1000410	0. 94
	MAMMA1000416	F-MAMMA1000416	0. 89
	MAMMA1000421	F-MAMMA1000421	0. 50
	MAMMA1000422	F-MAMMA1000422	0. 92
45	MAMMA1000429	F-MAMMA1000429	0. 92
	MAMMA1000458	F-MAMMA1000458	0.88
	MAMMA1000468	F-MAMMA1000468	0. 60
	MAMMA1000472	F-MAMMA1000472	0. 33
50	MAMMA1000490	F-MAMMA1000490	0.34
	MAMMA1000524	F-MAMMA1000524	0. 39
	MAMMA1000567	F-MAMMA1000567	0.40
	MAMMA1000583	F-MAMMA1000583	0. 45
55	MAMMA1000612	F-MAMMA1000612	0.89
	MAMMA1000623	F-MAMMA1000623	0. 44
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	MAMMA1000625	F-MAMMA1000625	0. 52
	MAMMA1000664	F-MAMMA1000664	0. 42
	MAMMA1000670	F-MAMMA1000670	0. 94
5	MAMMA1000672	F-MAMMA1000672	0. 44
	MAMMA1000684	F-MAMMA1000684	0.94
	MAMMA1000713	F-MAMMA1000713	0. 88
	MAMMA1000713		0. 67
10	MAMMA1000731		0. 40
10			
	MAMMA1000738		0. 93
	MAMMA1000746		0. 48
	MAMMA1000760		0. 94
15	MAMMA1000776	F-MAMMA1000776	0. 94
	MAMMA1000824	F-MAMMA1000824	0. 81
	MAMMA1000831	F-MAMMA1000831,	
	MAMMA1000841	F-MAMMA1000841	
20	MAMMA1000842	F-MAMMA1000842	0. 61
	MAMMA1000843	F-MAMMA1000843	0. 48
	MAMMA1000856	F-MAMMA1000856	0. 31
	MAMMA1000859	F-MAMMA1000859	0. 44
25	MAMMA1000865	F-MAMMA1000865	0. 94
	MAMMA1000875	F-MAMMA1000875	0. 49
	MAMMA1000897	F-MAMMA1000897	0. 94
	MAMMA1000906	F-MAMMA1000906	0. 45
30	MAMMA1000908	F-MAMMA1000908	0. 39
30	MAMMA1000914	F-MAMMA1000914	0.94
	MAMMA1000921	F-MAMMA1000921	0. 34
	MAMMA1000956	F-MAMMA1000956	0.39
05	MAMMA1000968	F-MAMMA1000968	0.35
35	MAMMA1000979	F-MAMMA1000979	0.68
	MAMMA1001008	F-MAMMA1001008	0.36
	MAMMA1001021	F-MAMMA1001021	0.48
	MAMMA1001041	F-MAMMA1001041	0, 34
40	MAMMA1001059	F-MAMMA1001059	0. 94
	MAMMA1001073	F-MAMMA1001073	0.94
	MAMMA1001075	F-MAMMA1001075	0. 94
	MAMMA1001078	F-MAMMA1001078	
45	MAMMA1001080	F-MAMMA1001080	0. 87
	MAMMA1001091	F-MAMMA1001091	0. 53
	MAMMA1001105	F-MAMMA1001105	0. 94
	MAMMA1001110	F-MAMMA1001110	0. 78
50	MAMMA1001126	F-MAMMA1001126	0. 78
	MAMMA1001120	F-MAMMA1001139	0. 38
	MAMMA1001139	F-MAMMA1001143	0. 79
	MAMMA1001154	F-MAMMA1001143	0.79
55	MAMMA1001181		
		F-MAMMA1001181	0.69
	MAMMA1001198	F-MAMMA1001198	0. 88

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	MAMMA1001202	F-MAMMA1001202	0.38
	MAMMA1001215	F-MAMMA1001215	0.94
	MAMMA1001222		0.89
5	MAMMA1001244	F-MAMMA1001244	
		F-MAMMA1001259	0. 91
		F-MAMMA1001260	0.75
	MAMMA1001271	F-MAMMA1001271	0.77
10		F-MAMMA1001291	0.77
			0. 75
		F-MAMMA1001296	
		F-MAMMA1001305	0.66
15		F-MAMMA1001343	0. 38
15		F-MAMMA1001346	0.81
		F-MAMMA1001388	0. 94
	MAMMA1001411	F-MAMMA1001411,	0. 42
	MAMMA1001419	F-MAMMA1001419	0. 45
20	MAMMA1001465	F-MAMMA1001465	0. 94
	MAMMA1001476	F-MAMMA1001476	0.90
	MAMMA1001487	F-MAMMA1001487	0. 78
	MAMMA1001510	F-MAMMA1001510	0. 82
25	MAMMA1001522	F-MAMMA1001522	0. 63
	MAMMA1001551	F-MAMMA1001551	0. 81
	MAMMA1001576	F-MAMMA1001576	0.94
	MAMMA1001600	F-MAMMA1001600	0. 72
30	MAMMA1001604	F-MAMMA1001604	0.87
	MAMMA1001620	F-MAMMA1001620	0. 43
	MAMMA1001627	F-MAMMA1001627	0.39
	MAMMA1001630	F-MAMMA1001630	0.46
35	MAMMA1001633	F-MAMMA1001633	0.58
	MAMMA1001635	F-MAMMA1001635	0.89
	MAMMA1001649	F-MAMMA1001649	0.62
	MAMMA1001654	F-MAMMA1001654	0.94
40	MAMMA1001692	F-MAMMA1001692	0. 83
40	MAMMA1001730	F-MAMMA1001730	0.48
	MAMMA1001735	F-MAMMA1001735	0.94
	MAMMA1001743	F-MAMMA1001743	0.94
45	MAMMA1001744	F-MANNA1001744	0. 33
45	MAMMA1001751	F-MAMMA1001751	0.94
	MAMMA1001754	F-MAMMA1001754	0. 47
	MAMMA1001757	F-MAMMA1001757	0. 38
	MAMMA1001764	F-MAMMA1001764	0.57
50	MAMMA1001768	F-MAMMA1001768	0. 85
	MAMMA1001771	F-MAMMA1001771	0. 39
	MAMMA1001783	F-MAMMA1001783	0. 61
	MAMMA1001785	F-MAMMA1001785	0. 41
55	MAMMA1001790	F-MAMMA1001790	0.69
	MAMMA1001812	F-MAMMA1001812	0. 32

	MAMMA1001817	F-MAMMA1001817	0. 55
	MAMMA1001824	F-MAMMA1001824	0. 57
	MAMMA1001837	F-MAMMA1001837	0.35
5	MAMMA1001848	F-MAMMA1001848	
	MAMMA1001851	F-MAMMA1001851	0. 46
	MAMMA1001858	F-MAMMA1001858	0. 53
	MAMMA1001864	F-MAMMA1001864	0. 32
10	MAMMA1001868		0.48
		F-MAMMA1001874	0, 94
	MAMMA1001956		0.47
	MAMMA1001969	F-MAMMA1001969	0. 31
15	MAMMA1001970	F-MAMMA1001970	0.34
	MAMMA1002009	F-MAMMA1002009	0. 45
	MAMMA1002011	F-MAMMA1002011,	
	MAMMA1002033	•	0. 33
20	MAMMA1002041		0.86
	MAMMA1002042		0.88
	MAMMA1002047		0.41
	MAMMA1002068		0. 38
25	MAMMA1002143		0. 91
	MAMMA1002153	F-MAMMA1002153	0.94
	MAMMA1002156	F-MAMMA1002156	0.73
	MAMMA1002170	F-MAMMA1002170	0. 43
30	MAMMA1002174	F-MAMMA1002174	0. 69
	MAMMA1002198	F-MAMMA1002198	0. 65
	MAMMA1002209	F-MAMMA1002209	0.80
	MAMMA1002219	F-MAMMA1002219	0.85
35	MAMMA1002236	F-MAMMA1002236	0.55
	MAMMA1002243	F-MAMMA1002243	0.50
	MAMMA1002268	F-MAMMA1002268	0. 43
	MAMMA1002269	F-MAMMA1002269	0. 33
40	MAMMA1002292	F-MAMMA1002292	0.76
	MAMMA1002294	F-MAMMA1002294	0.94
	MAMMA1002297	F-MAMMA1002297	0.91
	MAMMA1002308.	F-MAMMA1002308	0. 68
45	MAMMA1002312	F-MAMMA1002312	0.77
40	MAMMA1002317	F-MAMMA1002317	0.46
	MAMMA1002319	F-MAMMA1002319	0.31
	MAMMA1002329	F-MAMMA1002329	0.34
50	MAMMA1002333	F-MAMMA1002333	0.65
50	MAMMA1002351	F-MAMMA1002351	0.89
	MAMMA1002353	F-MAMMA1002353	0.94
	MAMMA1002355	F-MAMMA1002355	0.55
E E	MAMMA1002356	F-MAMMA1002356	0. 32
55	MAMMA1002362	F-MAMMA1002362	0.83
	MAMMA1002380	F-MAMMA1002380	0. 94

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	MAMMA1002384	F-MAMMA1002384	0. 42
	MAMMA1002385	F-MAMMA1002385	0. 82
	MAMMA1002413	F-MAMMA1002413	0. 94
5	MAMMA1002427	F-MAMMA1002427	0.40
	MAMMA1002434	F-MAMMA1002434	0.74
	MAMMA1002454	F-MAMMA1002454	0. 82
	MAMMA1002461	F-MAMMA1002461	0.49
10	MAMMA1002470	F-MAMMA1002470	0. 89
	MAMMA1002485		0. 55
	MAMMA1002524		0. 33
	MAMMA1002530		0.75
15	MAMMA1002554		0. 41
	MAMMA1002585		0. 31
	MAMMA1002590		
	MAMMA1002598	F-MAMMA1002598	0. 72
20			
20	MAMMA1002617	F-MAMMA1002617 F-MAMMA1002619	0. 49
	MAMMA1002619		0. 88
	MAMMA1002637		0. 94
•	MAMMA1002650		0. 54
25	MAMMA1002655		0. 39
	MAMMA1002665	F-MAMMA1002665	0. 33
	MAMMA1002671		0. 94
	MAMMA1002673		0. 70
30	MAMMA1002684		0. 94
	MAMMA1002685		0.88
	MAMMA1002699		0. 94
	MAMMA1002711	F-MAMMA1002711	0.39
35	MAMMA1002769	F-MAMMA1002769	0.88
	MAMMA1002775	F-MAMMA1002775	0.81
	MAMMA1002782	F-MAMMA1002782	0.87
	MAMMA1002796	F-MAMMA1002796	0.40
40	MAMMA1002807	F-MAMMA1002807	0.64
	MAMMA1002842	F-MAMMA1002842	0.94
	MAMMA1002843	F-MAMMA1002843	0. 32
	MAMMA1002868	F-MAMMA1002868	0.54
15	MAMMA1002869	F-MAMMA1002869	0.94
45	MAMMA1002881	F-MAMMA1002881	0.94
	MAMMA1002886	F-MAMMA1002886	0. 61
	MAMMA1002890	F-MAMMA1002890	0. 94
	MAMMA1002895	F-MAMMA1002895	0. 94
50	MAMMA1002937	F-MAMMA1002937	0. 91
	MAMMA1002938	F-MAMMA1002938	0. 78
	MAMMA1002947	F-MAMMA1002947	0. 94
	MAMMA1002964	F-MAMMA1002964	0. 34
55	MAMMA1002987	F-MAMMA1002987	0. 74
	MAMMA1003011	F-MAMMA1003011	0. 90
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	MAMMA1003013	F-MAMMA1003013	0.36
	MAMMA1003015	F-MAMMA1003015	0, 92
	MAMMA1003035	F-MAMMA1003035	0.94
5	MAMMA1003044	F-MAMMA1003044	
	MAMMA1003047	F-MAMMA1003047	0. 94
	MAMMA1003049	F-MAMMA1003049	0. 31
	MAMMA1003056	F-MAMMA1003056	0. 75
10	MAMMA1003057	F-MAMMA1003057	0. 33
	MAMMA1003066	F-MAMMA1003066	0. 38
	MAMMA1003099	F-MAMMA1003099	0. 79
•	MAMMA1003104	F-MAMMA1003104	0. 44
15	MAMMA1003113	F-MAMMA1003113	0. 90
	MAMMA1003127	F-MAMMA1003127	0. 50
	MAMMA1003127	F-MAMMA1003127	
	MAMMA1003135	•	
20	MAMMA1003150	F-MAMMA1003146	0.88
20	MAMMA1003166	F-MAMMA1003150	0. 73
	NT2RM1000001	F-MAMMA1003166	0. 94
		F-NT2RM1000001	0. 66
25	NT2RM1000018 NT2RM1000032	F-NT2RM1000018	0. 55
25	NT2RM1000032	F-NT2RM1000032 F-NT2RM1000035	0. 74
	NT2RM1000035		0. 52
	NT2RM1000037	F-NT2RM1000037	0. 94
	NT2RM1000039	F-NT2RM1000039	0.77
30	NT2RM1000059	F-NT2RM1000055	0.89
	NT2RM1000059	F-NT2RM1000059	0.41
	NT2RM1000082	F-NT2RM1000062	0. 65
	NT2RM1000080	F-NT2RM1000080	0.61
35	NT2RM1000086		0.94
	NT2RM1000092	F-NT2RM1000092 F-NT2RM1000118	0.94
	NT2RM1000118		0.76
	NT2RM1000119	F-NT2RM1000119 F-NT2RM1000127	0. 41
40	NT2RM1000127		0.94
	NT2RM1000131	F-NT2RM1000131 F-NT2RM1000132	0.90
	NT2RM1000153	F-NT2RM1000132	0.94
	NT2RM1000186		0. 90
45	NT2RM1000187	F-NT2RM1000186	0. 78
	NT2RM1000187	F-NT2RM1000187	0.54
		F-NT2RM1000199	0.33
	NT2RM1000242	F-NT2RM1000242	0.78
50	NT2RM1000244	F-NT2RM1000244	0.31
	NT2RM1000252	F-NT2RM1000252	0.76
	NT2RM1000256	F-NT2RM1000256	0.94
	NT2RH1000257	F-NT2RM1000257	0.89
55	NT2RM1000260	F-NT2RM1000260	0. 42
	NT2RM1000271	F-NT2RM1000271	0.74
	NT2RM1000272	F-NT2RM1000272	0. 78

	NT2RM1000280	F-NT2RM1000280	0. 94
	NT2RM1000300	F-NT2RM1000300	0.47
	NT2RM1000314	F-NT2RM1000314	0.61
5	NT2RM1000314	F-NT2RM1000318	
	NT2RM1000318	F-NT2RM1000318	0.70
	NT2RM1000341		0. 70
		F-NT2RM1000354	
40	NT2RM1000355	F-NT2RM1000355	0.94
10	NT2RM1000365	F-NT2RM1000365	0.31
	NT2RM1000377	F-NT2RM1000377	0.87
	NT2RM1000388	F-NT2RM1000388	0.94
	NT2RM1000394	F-NT2RM1000394	0. 58
15	NT2RM1000399	F-NT2RM1000399	0.86
	NT2RM1000421	F-NT2RM1000421	0. 39
	NT2RM1000430	F-NT2RM1000430,	
	NT2RM1000499	F-NT2RM1000499	0. 44
20	NT2RM1000539	F-NT2RM1000539	0. 87
	NT2RM1000553	F-NT2RM1000553	0. 88
	NT2RM1000555	F-NT2RM1000555	0. 37
	NT2RM1000563	F-NT2RM1000563	0. 63
25	NT2RM1000623	F-NT2RM1000623	0. 34
	NT2RM1000648	F-NT2RM1000648	0.94
	NT2RM1000661	F-NT2RM1000661	0.61
	NT2RM1000666	F-NT2RM1000666	0.94
30	NT2RM1000669	F-NT2RM1000669	0.94
	NT2RM1000672	F-NT2RM1000672	0. 53
	NT2RM1000691	F-NT2RM1000691	0.38
	NT2RM1000699	F-NT2RM1000699	0. 32
35	NT2RM1000702	F-NT2RM1000702	0. 74
30	NT2RM1000725	F-NT2RM1000725	0. 75
	NT2RM1000741	F-NT2RM1000741	0. 94
	NT2RM1000742	F-NT2RM1000742	0. 54
40	NT2RM1000746	F-NT2RM1000746	0. 94
40	NT2RM1000770	F-NT2RM1000770	0. 48
	NT2RM1000772	F-NT2RM1000772	0. 39
	NT2RM1000780	F-NT2RM1000780	0. 66
	NT2RM1000781	F-NT2RM1000781	0.49
45	NT2RM1000800	F-NT2RM1000800	0. 72
	NT2RM1000802	F-NT2RM1000802	0.89
	NT2RM1000811	F-NT2RM1000811	0. 53
	NT2RM1000826	F-NT2RM1000826	0. 39
50	NT2RM1000829	F-NT2RM1000829	0. 63
	NT2RM1000833	F-NT2RM1000833	0.94
	NT2RM1000850	F-NT2RM1000850	0.89
	NT2RM1000852	F-NT2RM1000852	0.88
55	NT2RM1000857	F-NT2RM1000857	0. 58
	NT2RM1000867	F-NT2RM1000867	0.90

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	NT2RM1000874	F-NT2RM1000874	0. 89
	NT2RM1000882	F-NT2RM1000882	0. 63
	NT2RM1000883	F-NT2RM1000883	0.71
5	NT2RM1000885	F-NT2RM1000885	0. 89
	NT2RM1000894	F-NT2RM1000894	0. 57
	NT2RM1000898	F-NT2RM1000898	0. 87
	NT2RM1000905	F-NT2RM1000905	0. 94
10	NT2RM1000924	F-NT2RM1000924	0.88
	NT2RM1000927		0. 70
	NT2RM1000962	F-NT2RM1000962	0. 84
	NT2RM1000978	F-NT2RM1000978	0. 39
15	NT2RM1001003	F-NT2RM1001003	0. 81
	NT2RM1001008	F-NT2RM1001008	0. 94
	NT2RM1001043	F-NT2RM1001043	1
	NT2RM1001044	F-NT2RM1001044	0. 94
20	NT2RM1001059	F-NT2RM1001059	0. 94
	NT2RM1001066		0. 94
	NT2RM1001072	F-NT2RM1001072	0.77
,	NT2RM1001074	F-NT2RM1001074	0. 94
25	NT2RM1001082	F-NT2RM1001082	0. 94
	NT2RM1001085	F-NT2RM1001085	0. 83
	NT2RM1001092	F-NT2RM1001092	0.77
	NT2RM1001102	F-NT2RM1001102	0. 59
30	NT2RM1001105	F-NT2RM1001105	0. 94
	NT2RM1001112	F-NT2RM1001112	0. 50
	NT2RM1001115	F-NT2RM1001115	0.32
	NT2RM1001139	F-NT2RM1001139	0. 77
35	NT2RM2000006	F-NT2RM2000006	0.34
	NT2RM2000013	F-NT2RM2000013	0.70
	NT2RM2000030	F-NT2RM2000030	0.89
	NT2RM2000032	F-NT2RM2000032	0. 36
40	NT2RM2000042	F-NT2RM2000042	0. 32
	NT2RM2000092	F-NT2RM2000092	0. 55
	NT2RM2000093	F-NT2RM2000093	0. 33
	NT2RM2000101	F-NT2RM2000101	0. 77
45	NT2RM2000124	F-NT2RM2000124	0. 94
	NT2RM2000191	F-NT2RM2000191	0.80
	NT2RM2000192	F-NT2RM2000192	0. 93
	NT2RM2000239	F-NT2RM2000239	0. 68
50	NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN	F-NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN	0.94
	NT2RM2000250	F-NT2RM2000250	0.94
	NT2RM2000259	F-NT2RM2000259 F-NT2RM2000260	0. 68 0. 37
	NT2RM2000287	F-NT2RM2000287	0. 37
55	NT2RM2000322	F-NT2RM2000322	0. 46
	NT2RM2000322	F-NT2RM2000359	0. 94
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	NT2RM2000363	F-NT2RM2000363	0.91
	NT2RM2000368	F-NT2RM2000368	0. 34
	NT2RM2000371	F-NT2RM2000371	0.94
5	NT2RM2000374	F-NT2RM2000374	
	NT2RM2000395	F-NT2RM2000395	0. 33
	NT2RM2000402	F-NT2RM2000402	0.80
•	NT2RM2000407	F-NT2RM2000407	0.88
10	NT2RM2000420	F-NT2RM2000420	0. 92
	NT2RM2000422	F-NT2RM2000422	0.49
	NT2RM2000452	F-NT2RM2000452	0. 93
	NT2RM2000469	F-NT2RM2000469	0.39
15	NT2RM2000409	F-NT2RM2000490	0.90
	NT2RM2000490	F-NT2RM2000502	0. 64
	NT2RM2000504	F-NT2RM2000504	
	NT2RM2000504	F-NT2RM2000522	
20	NT2RM2000540	F-NT2RM2000540	0. 91
	NT2RM2000540		1
	NT2RM2000556	F-NT2RM2000556	0.91
		F-NT2RM2000566	0.39
25	NT2RM2000567	F-NT2RM2000567	0. 44
23	NT2RM2000569	F-NT2RM2000569	0. 49
	NT2RM2000577	F-NT2RM2000577	0. 45
	NT2RM2000581	F-NT2RM2000581	0. 72
••	NT2RM2000588	F-NT2RM2000588	0.31
30	NT2RM2000594	F-NT2RM2000594	0.79
	NT2RM2000599	F-NT2RM2000599	0.80
	NT2RM2000612	F-NT2RM2000612	0. 75
	NT2RM2000623	F-NT2RM2000623	0.72
35	NT2RM2000624	F-NT2RM2000624	0. 94
	NT2RM2000635	F-NT2RM2000635	0. 73
	NT2RM2000636	F-NT2RM2000636	0. 45
	NT2RM2000639	F-NT2RM2000639	0.43
40	NT2RM2000649	F-NT2RM2000649	0.74
	NT2RM2000669	F-NT2RM2000669	0. 94
	NT2RM2000691	F-NT2RM2000691	0.84
	NT2RM2000714	F-NT2RM2000714	0. 92
45	NT2RM2000718	F-NT2RM2000718	0. 52
	NT2RM2000735	F-NT2RM2000735	0. 33
	NT2RM2000740	F-NT2RM2000740	0. 94
	NT2RM2000795	F-NT2RM2000795	0. 44
50	NT2RM2000821	F-NT2RM2000821	0. 33
	NT2RM2000837	F-NT2RM2000837	0.57
	NT2RM2000951	F-NT2RM2000951	0.32
	NT2RM2000952	F-NT2RM2000952	0.50
55	NT2RM2000984	F-NT2RM2000984	0. 76
	NT2RM2001004	F-NT2RM2001004	0. 55
	NT2RM2001035	F-NT2RM2001035	0. 82

	NT2RM2001065	F-NT2RM2001065	0.65
	NT2RM2001100	F-NT2RM2001100	0.89
_	NT2RM2001105	F-NT2RM2001105	0.94
5	NT2RM2001131	F-NT2RM2001131_	0.94
	NT2RM2001141	F-NT2RM2001141	0.74
	NT2RM2001152	F-NT2RM2001152	0.33
	NT2RM2001177	F-NT2RM2001177	0.32
10	NT2RM2001194	F-NT2RM2001194	0. 83
	NT2RM2001196	F-NT2RM2001196	0. 42
	NT2RM2001201	F-NT2RM2001201	0.90
	NT2RM2001221	F-NT2RM2001221	0.39
15	NT2RM2001238	F-NT2RM2001238	0.94
	NT2RM2001243	F-NT2RM2001243	0.44
	NT2RM2001247	F-NT2RM2001247,	0.42
	NT2RM2001256	F-NT2RM2001256	0.60
20	NT2RM2001291	F-NT2RM2001291	0.39
	NT2RM2001306	F-NT2RM2001306	0.81
	NT2RM2001312	F-NT2RM2001312	0.64
	NT2RM2001319	F-NT2RM2001319	0.53
25	NT2RM2001324	F-NT2RM2001324	0.94
	NT2RM2001345	F-NT2RM2001345	0. 94
	NT2RM2001360	F-NT2RM2001360	0.69
	NT2RM2001370		0. 44
30	NT2RM2001393	F-NT2RM2001393	0. 94
	NT2RM2001420	F-NT2RM2001420	0. 46
	NT2RM2001424		0. 94
	NT2RM2001499	F-NT2RM2001499	0.63
35	NT2RM2001504	F-NT2RM2001504	0. 94
	NT2RM2001524	F-NT2RM2001524	0.76
	NT2RM2001544		0. 59
	NT2RM2001547		0.48
40	NT2RM2001575	F-NT2RM2001575	0. 91
	NT2RM2001582	F-NT2RM2001582	0.34
	NT2RM2001588	F-NT2RM2001588	0.48
	NT2RM2001592 NT2RM2001605	F-NT2RM2001592 F-NT2RM2001605	0. 64 0. 47
45	NT2RM2001605	•	0. 61
	NT2RM2001613	F-NT2RM2001613	0. 31
		F-NT2RM2001635	
	NT2RM2001635 NT2RM2001637		0. 39
50	NT2RM2001637	F-NT2RM2001637	0. 71
	NT2RM2001641		0.73
	NT2RM2001652		0.40
	NT2RM2001659		0. 40
55	NT2RM2001664		0.53
	NT2RM2001668	•	0.64
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	NT2DN2001670	E NT00000004070	0.50
	NT2RM2001670	F-NT2RM2001670	0.59
	NT2RM2001671	F-NT2RM2001671	0.41
	NT2RM2001675	F-NT2RM2001675	0.51
5	NT2RM2001681	F-NT2RM2001681	
	NT2RM2001688	F-NT2RM2001688	0. 74
	NT2RM2001695	F-NT2RM2001695	0.31
	NT2RM2001696	F-NT2RM2001696	0.94
10	NT2RM2001698	F-NT2RM2001698	0. 69
	NT2RM2001699	F-NT2RM2001699	0. 85
	NT2RM2001700	F-NT2RM2001700	0.71
	NT2RM2001706	F-NT2RM2001706	0.38
15	NT2RM2001716	F-NT2RM2001716	0. 78
	NT2RM2001718	F-NT2RM2001718	0.81
	NT2RM2001723	F-NT2RM2001723,	0.83
	NT2RM2001727	F-NT2RM2001727	0. 87
20	NT2RM2001730	F-NT2RM2001730	0. 94
	NT2RM2001743	F-NT2RM2001743	0, 94
	NT2RM2001753	F-NT2RM2001753	0. 75
	NT2RM2001760	F-NT2RM2001760	0. 75
25	NT2RM2001768	F-NT2RM2001768	0. 71
20	NT2RM2001771	F-NT2RM2001771	0. 33
	NT2RM2001782	F-NT2RM2001782	0.34
	NT2RM2001784	F-NT2RM2001784	0. 45
30	NT2RM2001785	F-NT2RM2001785	0. 94
30	NT2RM2001797	F-NT2RM2001797	0.74
	NT2RM2001800	F-NT2RM2001800	0.90
	NT2RM2001803	F-NT2RM2001803	0.36
0.5	NT2RM2001805	F-NT2RM2001805	0. 81
35	NT2RM2001813	F-NT2RM2001813	0.94
	NT2RM2001823	F-NT2RM2001823	0. 94
	NT2RM2001839	F-NT2RM2001839	0.39
	NT2RM2001840	F-NT2RM2001840	0. 94
40	NT2RM2001855	F-NT2RM2001855	0.92
	NT2RM2001867	F-NT2RM2001867	0.82
	NT2RM2001879	F-NT2RM2001879	0.86
	NT2RM2001886	F-NT2RM2001886	0.91
45	NT2RM2001896	F-NT2RM2001896	0. 94
	NT2RM2001903	F-NT2RM2001903	0. 61
	NT2RM2001930	F-NT2RM2001930	0. 94
	NT2RM2001935	F-NT2RM2001935	0. 77
50	NT2RM2001936	F-NT2RM2001936	0. 94
	NT2RM2001950	F-NT2RM2001950	0.34
	NT2RM2001982	F-NT2RM2001982	0.48
	NT2RM2001983	F-NT2RM2001983	0.59
55	NT2RM2001989	F-NT2RM2001989	0.94
	NT2RM2001997	F-NT2RM2001997	0. 90

	NT2RM2001998	F-NT2RM2001998	0, 72
	NT2RM2002004	F-NT2RM2002004	0. 46
	NT2RM2002014	F-NT2RM2002014	0. 32
5	NT2RM2002030	F-NT2RM2002030	i
	NT2RM2002049	F-NT2RM2002049	0, 59
	NT2RM2002055	F-NT2RM2002055	0.94
	NT2RM2002088	F-NT2RM2002088	0.47
10	NT2RM2002091	F-NT2RM2002091	0. 78
•	NT2RM2002100	F-NT2RM2002100	0. 87
	NT2RM2002109	F-NT2RM2002109	0. 86
	NT2RM2002128	F-NT2RM2002128	0. 42
15	NT2RM2002142	F-NT2RM2002142	0. 94
	NT2RM2002145	F-NT2RM2002145	0. 94
	NT2RM2002178	F-NT2RM2002178	
	NT2RM2002580	F-NT2RM2002580	0. 94
20	NT2RM4000024	F-NT2RM4000024	0.76
	NT2RM4000027	F-NT2RM4000027	0. 40
	NT2RM4000030	F-NT2RM4000030	0. 66
	NT2RM4000061	F-NT2RM4000061	0. 33
25	NT2RM4000104	F-NT2RM4000104	0. 93
	NT2RM4000139	F-NT2RM4000139	0.87
	NT2RM4000155	F-NT2RM4000155	0.51
	NT2RM4000156	F-NT2RM4000156	0.39
30	NT2RM4000167	F-NT2RM4000167	0.50
	NT2RM4000169	F-NT2RM4000169	0.44
	NT2RM4000191	F-NT2RM4000191	0.94
	NT2RM4000197	F-NT2RM4000197	0.41
35	NT2RM4000199	F-NT2RM4000199	0.76
	NT2RM4000202	F-NT2RM4000202	0.47
	NT2RM4000210	F-NT2RM4000210	0.44
	NT2RM4000215	F-NT2RM4000215	0.94
40	NT2RM4000229	F-NT2RM4000229	0. 93
	NT2RM4000233	F-NT2RM4000233	0. 74
,	NT2RM4000251	F-NT2RM4000251	0. 48
	NT2RM4000290	F-NT2RM4000290	0.94
45	NT2RM4000324	F-NT2RM4000324	0. 32
	NT2RM4000344	F-NT2RM4000344	0.94
	NT2RM4000349	F-NT2RM4000349	0. 78
	NT2RM4000354	F-NT2RM4000354	0. 36
50	NT2RM4000356	F-NT2RM4000356	0. 58
	NT2RM4000386	F-NT2RM4000386	0. 90
	NT2RM4000395	F-NT2RM4000395	0. 47
	NT2RM4000421	F-NT2RM4000421	0. 46
55	NT2RM4000433	F-NT2RM4000433	0. 48
33	NT2RM4000457	F-NT2RM4000457	0. 75
	NT2RM4000471	F-NT2RM4000471	0. 92

	NT2RM4000486	F-NT2RM4000486	0. 55
	NT2RM4000496	F-NT2RM4000486	0. 62
	NT2RM4000511	F-NT2RM4000498	0. 62
5	NT2RM4000514	F-NT2RM4000514	
	NT2RM4000515	F-NT2RM4000515	0.66
	NT2RM4000520	F-NT2RM4000520	0.69
10	NT2RM4000531	F-NT2RM4000531	0, 62
	NT2RM4000534	F-NT2RM4000534	0.94
	NT2RM4000585	F-NT2RM4000585	0. 52
	NT2RM4000590	F-NT2RM4000590	0, 94
45	NT2RM4000595	F-NT2RM4000595	0, 44
15	NT2RM4000603	F-NT2RM4000603	0. 94
	NT2RM4000611	F-NT2RM4000611	0. 46
	NT2RM4000616	F-NT2RM4000616,	0. 94
	NT2RM4000674	F-NT2RM4000674	0.80
20	NT2RM4000689	F-NT2RM4000689	0. 58
	NT2RM4000698	F-NT2RM4000698	0.71
	NT2RM4000700	F-NT2RM4000700	0.59
	NT2RM4000712	F-NT2RM4000712	0.94
25	NT2RM4000717	F-NT2RM4000717	0.69
	NT2RM4000733	F-NT2RM4000733	0.94
	NT2RM4000734	F-NT2RM4000734	0.42
	NT2RM4000741	F-NT2RM4000741	0.83
30	NT2RM4000751	F-NT2RM4000751	0.91
	NT2RM4000764	F-NT2RM4000764	0.35
	NT2RM4000778	F-NT2RM4000778	0.90
	NT2RM4000787	F-NT2RM4000787	0.40
35	NT2RM4000790	F-NT2RM4000790	0.62
	NT2RM4000795	F-NT2RM4000795	0.94
	NT2RM4000796	F-NT2RM4000796	0.93
	NT2RM4000798	F-NT2RM4000798	0.61
40	NT2RM4000813	F-NT2RM4000813	0.58
10	NT2RM4000820	F-NT2RM4000820	0.39
	NT2RM4000833	F-NT2RM4000833	0.80
	NT2RM4000848	F-NT2RM4000848	0.73
45	NT2RM4000852	F-NT2RM4000852	0.74
45	NT2RM4000887	F-NT2RM4000887	0.42
	NT2RM4000895	F-NT2RM4000895	0. 47
	NT2RM4000950	F-NT2RM4000950	0.53
	NT2RM4000971	F-NT2RM4000971	0.73
50	NT2RM4000996	F-NT2RM4000996	0.42
	NT2RM4001002	F-NT2RM4001002	0. 57
	NT2RM4001032	F-NT2RM4001032	0.94
	NT2RM4001047	F-NT2RM4001047	0.76
55	NT2RM4001054	F-NT2RM4001054	0.91
	NT2RM4001084	F-NT2RM4001084	0.94

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	NT2RM4001092	F-NT2RM4001092	0.94
	NT2RM4001116	F-NT2RM4001116	0.36
	NT2RM4001140	F-NT2RM4001140	0.82
5	NT2RM4001151	F-NT2RM4001151	
•	NT2RM4001155	F-NT2RM4001155	0.94
	NT2RM4001187	F-NT2RM4001187	0.61
	NT2RM4001191	F-NT2RM4001191	0. 77
40		F-NT2RM4001200	0.77
10	NT2RM4001200		0. 72
	NT2RM4001203	F-NT2RM4001203	
	NT2RM4001204	F-NT2RM4001204	0.89
	NT2RM4001217	F-NT2RM4001217	0.93
15	NT2RM4001256	F-NT2RM4001256	0.89
	NT2RM4001258	F-NT2RM4001258	0.94
	NT2RM4001309	F-NT2RM4001309	
	NT2RM4001316	F-NT2RM4001316	0.38
20	NT2RM4001320	F-NT2RM4001320	0.33
	NT2RM4001340	F-NT2RM4001340	0.83
•	NT2RM4001344	F-NT2RM4001344	0.94
	NT2RM4001347	F-NT2RM4001347	0.88
25	NT2RM4001371	F-NT2RM4001371	0.61
	NT2RM4001382	F-NT2RM4001382	0. 93
	NT2RM4001384	F-NT2RM4001384	0.32
	NT2RM4001410	F-NT2RM4001410	0.94
30	NT2RM4001411	F-NT2RM4001411	0. 47
30	NT2RM4001412	F-NT2RM4001412	0. 85
	NT2RM4001444	F-NT2RM4001444	0.94
	NT2RM4001454	F-NT2RM4001454	0.72
05	NT2RM4001455	F-NT2RM4001455	0.39
35	NT2RM4001483	F-NT2RM4001483	0.76
	NT2RM4001489	F-NT2RM4001489	0.94
	NT2RM4001522	F-NT2RM4001522	0.88
	NT2RM4001565	F-NT2RM4001565	0. 78
40	NT2RM4001566	F-NT2RM4001566	0.87
	NT2RM4001569	F-NT2RM4001569	0.70
	NT2RM4001582	F-NT2RM4001582	0.76
	NT2RM4001592	F-NT2RM4001592	0.31
45	NT2RM4001594	F-NT2RM4001594	0.86
	NT2RM4001597	F-NT2RM4001597	0. 91
	NT2RM4001611	F-NT2RM4001611	0. 91
	NT2RM4001629	F-NT2RM4001629	0.72
50	NT2RM4001650	F-NT2RM4001650	0. 37
	NT2RM4001662	F-NT2RM4001662	0. 70
	NT2RM4001666	F-NT2RM4001666	0. 70
	NT2RM4001682	F-NT2RM4001682	0. 91
55	NT2RM4001710	F-NT2RM4001710	0. 45
55	NT2RM4001714	F-NT2RM4001714	0. 45
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		101701174	
	NT2RM4001715	F-NT2RM4001715	0. 33
	NT2RM4001731	F-NT2RM4001731	0. 94
	NT2RM4001741		0. 87
5	NT2RM4001746	F-NT2RM4001746	i
	NT2RM4001758		0. 59
	NT2RM4001783	F-NT2RM4001783	0. 68
	NT2RM4001813		0. 39
10	NT2RM4001819		0. 43
70	NT2RM4001823		0. 94
	NT2RM4001828		0.83
	NT2RM4001836		0.61
45	NT2RM4001836		
15			0.37
	NT2RM4001842	F-NT2RM4001842	0.68
	NT2RM4001865	,	- 1
	NT2RM4001876	F-NT2RM4001876	0.88
20	NT2RM4001880		0. 43
	NT2RM4001905		0. 94
	NT2RM4001922		0.48
	NT2RM4001938		0. 67
25	NT2RM4001940		0. 92
	NT2RM4001965		0. 94
	NT2RM4001969		0. 88
	NT2RM4001979		0. 42
30			0. 86
			0. 94
			0. 92
			0. 43
35			0. 42
			0.88
	NT2RM4002055		0. 32
			0.87
40	NT2RM4002063		0.49
10	NT2RM4002066		0.39
	NT2RM4002073		0.78
	NT2RM4002075		0.87
45	NT2RM4002093	F-NT2RM4002093	0.82
45	NT2RM4002109		0. 52
	NT2RM4002128		0. 68
	NT2RM4002140	F-NT2RM4002140	0.94
	NT2RM4002145	`	0. 94
50	NT2RM4002146	F-NT2RM4002146	0. 89
	NT2RM4002161	F-NT2RM4002161	0. 49
	NT2RM4002174		0. 36
	NT2RM4002189	F-NT2RM4002189	0. 34
55	NT2RM4002194	F-NT2RM4002194	0. 73
	NT2RM4002205	F-NT2RM4002205	0. 94

	NT2RM4002213	F-NT2RM4002213	0. 52
	NT2RM4002226	F-NT2RM4002226	0. 48
	NT2RM4002251	F-NT2RM4002251	0. 94
5	NT2RM4002256	F-NT2RM4002256_	
	NT2RM4002266	F-NT2RM4002266	0.73
	NT2RM4002294	F-NT2RM4002294	0.86
	NT2RM4002301	F-NT2RM4002301	0.40
10	NT2RM4002323	F-NT2RM4002323	0. 81
	NT2RM4002339	F-NT2RM4002339	0. 32
	NT2RM4002344	F-NT2RM4002344	0. 87
	NT2RM4002373	F-NT2RM4002373	0. 41
15	NT2RM4002374	F-NT2RM4002374	0. 33
	NT2RM4002398	F-NT2RM4002398	0. 94
	NT2RM4002409	F-NT2RM4002409	
	NT2RM4002446	F-NT2RM4002446	0. 61
20	NT2RM4002452	F-NT2RM4002452	0. 47
	NT2RM4002457	F-NT2RM4002457	0. 92
	NT2RM4002460	F-NT2RM4002460	0.61
	NT2RM4002482	F-NT2RM4002482	0. 94
25	NT2RM4002493	F-NT2RM4002493	0. 39
	NT2RM4002527	F-NT2RM4002527	0. 64
	NT2RM4002534	F-NT2RM4002534	0.76
	NT2RM4002558	F-NT2RM4002558	0. 62
30	NT2RM4002565	F-NT2RM4002565	0. 54
	NT2RM4002571	F-NT2RM4002571	0. 90
	NT2RM4002593	F-NT2RM4002593	0. 91
	NT2RM4002594	F-NT2RM4002594	0.70
35	NT2RM4002623	F-NT2RM4002623	0. 81
	NT2RP1000018	F-NT2RP1000018	0.31
	NT2RP1000035	F-NT2RP1000035	0. 49
	NT2RP1000040	F-NT2RP1000040	0. 45
40	NT2RP1000063	F-NT2RP1000063	0. 90
40	NT2RP1000086	F-NT2RP1000086	0.89
	NT2RP1000101	F-NT2RP1000101	0. 76
	NT2RP1000111	F-NT2RP1000111	0. 94
45	NT2RP1000112	F-NT2RP1000112	0. 45
45	NT2RP1000124	F-NT2RP1000124	0. 94
	NT2RP1000130	F-NT2RP1000130	0. 69
	NT2RP1000163	F-NT2RP1000163	0. 90
50	NT2RP1000170	F-NT2RP1000170	0. 94
30	NT2RP1000174	F-NT2RP1000174	0. 74
	NT2RP1000191	F-NT2RP1000191	0. 76
	NT2RP1000202	F-NT2RP1000202	0.60
55	NT2RP1000243	F-NT2RP1000243	0.81
55	NT2RP1000259	F-NT2RP1000259	0. 79
	NT2RP1000272	F-NT2RP1000272	0. 94

	NT2RP1000324	F-NT2RP1000324	0. 32
·	NT2RP1000326	F-NT2RP1000326	0. 94
	NT2RP1000333	F-NT2RP1000333	0.94
5	NT2RP1000348	F-NT2RP1000348_	0. 94
	NT2RP1000357	F-NT2RP1000357	0.87
	NT2RP1000358	F-NT2RP1000358	0.79
	NT2RP1000363	F-NT2RP1000363	0.40
10	NT2RP1000376	F-NT2RP1000376	0. 61
	NT2RP1000409	F-NT2RP1000409	0.85
	NT2RP1000413	F-NT2RP1000413	0. 42
	NT2RP1000416	F-NT2RP1000416	0. 68
15	NT2RP1000418	F-NT2RP1000418	0.88
	NT2RP1000439	F-NT2RP1000439	0.69
	NT2RP1000443	F-NT2RP1000443	
	NT2RP1000460	F-NT2RP1000440	0.86
20	NT2RP1000470	F-NT2RP1000470	0.56
20	NT2RP1000478	F-NT2RP1000478	0. 87
	NT2RP1000478	F-NT2RP1000478	0.58
	NT2RP1000481	F-NT2RP1000493	0.34
25	NT2RP1000513	F-NT2RP1000513	0.89
25	NT2RP1000522	F-NT2RP1000522	0. 34
	NT2RP1000547	F-NT2RP1000547	0. 89
	NT2RP1000574	F-NT2RP1000574	0. 74
	NT2RP1000577	F-NT2RP1000577	0. 53
30	NT2RP1000577	F-NT2RP1000577	0. 66
	NT2RP1000609	F-NT2RP1000609	0.90
	NT2RP1000629	F-NT2RP1000629	0. 94
	NT2RP1000630	F-NT2RP1000630	0. 93
35	NT2RP1000677	F-NT2RP1000677	0. 94
	NT2RP1000688	F-NT2RP1000688	0. 94
	NT2RP1000695	F-NT2RP1000695	0.89
	NT2RP1000701	F-NT2RP1000701	0. 89
40	NT2RP1000721	F-NT2RP1000721	0. 91
	NT2RP1000730	F-NT2RP1000730	0. 94
	NT2RP1000733	F-NT2RP1000733	0.90
	NT2RP1000738	F-NT2RP1000738	0.:64
45	NT2RP1000746	F-NT2RP1000746	0. 92
	NT2RP1000767	F-NT2RP1000767	0. 82
	NT2RP1000782	F-NT2RP1000782	0. 94
	NT2RP1000796	F-NT2RP1000796	0. 33
50	NT2RP1000825	F-NT2RP1000825	0.74
	NT2RP1000833	F-NT2RP1000833	0. 94
	NT2RP1000834	F-NT2RP1000834	0. 88
	NT2RP1000836	F-NT2RP1000836	0, 59
55	NT2RP1000846	F-NT2RP1000846	0. 46
	NT2RP1000851	F~NT2RP1000851	0. 82
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	NT2RP1000856	F-NT2RP1000856	0. 94
	NT2RP1000860	F-NT2RP1000860	0. 60
_	NT2RP1000902	F-NT2RP1000902	0. 42
5	NT2RP1000915	F-NT2RP1000915_	0, 90
	NT2RP1000916	F-NT2RP1000916	0. 67
	NT2RP1000943	F-NT2RP1000943	0. 77
	NT2RP1000944	F-NT2RP1000944	0. 94
10	NT2RP1000947	F-NT2RP1000947	0. 38
	NT2RP1000954	F~NT2RP1000954	0. 59
	NT2RP1000958	F-NT2RP1000958	0. 59
	NT2RP1000959	F-NT2RP1000959	0. 74
15	NT2RP1000966	F-NT2RP1000966	0. 93
	NT2RP1000980	F-NT2RP1000980	0.31
	NT2RP1000988	F-NT2RP1000988;	0. 53
	NT2RP1001011	F-NT2RP1001011	0. 65
20	NT2RP1001013	F-NT2RP1001013	0.34
	NT2RP1001014	F-NT2RP1001014	0.94
	NT2RP1001033	F-NT2RP1001033	0.40
	NT2RP1001073	F-NT2RP1001073	0.74
25	NT2RP1001079	F-NT2RP1001079	0.48
	NT2RP1001080	F-NT2RP1001080	0.91
	NT2RP1001113	F-NT2RP1001113	0. 94
	NT2RP1001173	F-NT2RP1001173	0.57
30	NT2RP1001177	F-NT2RP1001177	0.75
	NT2RP1001185	F-NT2RP1001185	0.94
	NT2RP1001199	F-NT2RP1001199	0. 42
	NT2RP1001247	F-NT2RP1001247	0. 67
35	NT2RP1001248	F-NT2RP1001248	0.89
	NT2RP1001253	F-NT2RP1001253	0.94
	NT2RP1001286	F-NT2RP1001286	0.59
	NT2RP1001294	F-NT2RP1001294	0.81
40	NT2RP1001302	F-NT2RP1001302	0.86
	NT2RP1001310	F-NT2RP1001310	0.91
	NT2RP1001311	F-NT2RP1001311	0.87
	NT2RP1001313 NT2RP1001361	F-NT2RP1001313	0.91
45	NT2RP1001385	F-NT2RP1001361	0.83
	NT2RP1001385	F-NT2RP1001385	0.36
	NT2RP1001393	F-NT2RP1001395 F-NT2RP1001410	0.84
			0.72
50	NT2RP1001424 NT2RP1001432	F-NT2RP1001424 F-NT2RP1001432	0. 81 0. 81
	NT2RP1001432	F-NT2RP1001449	0.81
		F-NT2RP1001449	0.77
	NT2RP1001467		0. 92
55	NT2RP1001466		0. 53
	NT2RP1001479		0. 34
		1 1001402	U. 04

	NT2RP1001494	F-NT2RP1001494	0. 93
	NT2RP1001543	F-NT2RP1001543	0. 94
	NT2RP1001546	F-NT2RP1001546	0. 94
5	NT2RP1001569		
	NT2RP1001616	F-NT2RP1001569_	0. 46
	NT2RP1001665	F-NT2RP1001665	0.94
10	NT2RP2000001	F-NT2RP2000001	0.94
•	NT2RP2000006	F-NT2RP2000006	0. 74
	NT2RP2000007	F-NT2RP2000007	0. 94
	NT2RP2000008	F-NT2RP2000008	0. 83
15	NT2RP2000032	F-NT2RP2000032	0. 94
70	NT2RP2000040	F-NT2RP2000040	0. 92
	NT2RP2000045	F-NT2RP2000045	0. 94
	NT2RP2000054	F-NT2RP2000054	
00	NT2RP2000056	F-NT2RP2000056	0. 42
20	NT2RP2000067	F-NT2RP2000067	0. 88
	NT2RP2000070	F-NT2RP2000070	0. 48
	NT2RP2000079	F-NT2RP2000079	0. 85
	NT2RP2000088	F-NT2RP2000088	0. 94
25	NT2RP2000091	F-NT2RP2000091	0. 79
	NT2RP2000097	F-NT2RP2000097	0.66
	NT2RP2000114	F-NT2RP2000114	0. 86
	NT2RP2000120	F-NT2RP2000120	0. 48
30	NT2RP2000126	F-NT2RP2000126	0. 77
	NT2RP2000133	F-NT2RP2000133	0. 64
	NT2RP2000147	F-NT2RP2000147	0. 94
	NT2RP2000153	F-NT2RP2000153	0. 87
35	NT2RP2000157	F-NT2RP2000157	0. 48
	NT2RP2000161	F-NT2RP2000161	0. 79
	NT2RP2000173	F-NT2RP2000173	0. 75
	NT2RP2000175	F-NT2RP2000175	0.79
40	NT2RP2000195	F-NT2RP2000195	0. 91
	NT2RP2000205	F-NT2RP2000205	0, 94
	NT2RP2000208	F-NT2RP2000208	0, 84
	NT2RP2000224	F-NT2RP2000224	0. 52
45	NT2RP2000232	F-NT2RP2000232	0. 33
	NT2RP2000233	F-NT2RP2000233	0. 57
	NT2RP2000239	F-NT2RP2000239	0. 52
	NT2RP2000248	F-NT2RP2000248	0.36
50	NT2RP2000274	FNT2RP2000274	0. 55
	NT2RP2000283	F-NT2RP2000283	0. 61
	NT2RP2000288	F-NT2RP2000288	0. 31
	NT2RP2000297	F-NT2RP2000297	0.78
55	NT2RP2000298	F-NT2RP2000298	0. 94
	NT2RP2000310	F-NT2RP2000310	0. 52
	NT2RP2000328	F-NT2RP2000328	0. 75

	NT2RP2000329	F-NT2RP2000329	0. 94
	NT2RP2000346	F-NT2RP2000346	0. 62
	NT2RP2000369	F-NT2RP2000369	0. 87
5	NT2RP2000412	F-NT2RP2000412	
	NT2RP2000414	F-NT2RP2000414	0.79
	NT2RP2000422	F-NT2RP2000422	0.46
	NT2RP2000438	F-NT2RP2000438	0. 94
10	NT2RP2000448	F-NT2RP2000448	0. 94
	NT2RP2000503	F-NT2RP2000503	0. 94
	NT2RP2000510	F-NT2RP2000510	0. 87
	NT2RP2000516	F-NT2RP2000516	0. 42
15	NT2RP2000617	F-NT2RP2000617	0. 94
	NT2RP2000634	F-NT2RP2000634	0.74
	NT2RP2000656	F-NT2RP2000656,	
	NT2RP2000658	F-NT2RP2000658	0. 55
20	NT2RP2000668	F-NT2RP2000668	0. 81
	NT2RP2000704	F-NT2RP2000704	0. 88
	NT2RP2000710	F-NT2RP2000710	0. 58
	NT2RP2000764	F-NT2RP2000764	0. 59
25	NT2RP2000809	F-NT2RP2000809	0.51
	NT2RP2000812	F-NT2RP2000812	0.37
	NT2RP2000814	F-NT2RP2000814	0.66
	NT2RP2000816	F-NT2RP2000816	0.50
30	NT2RP2000819	F-NT2RP2000819	0. 35
•	NT2RP2000841	F-NT2RP2000841	0. 93
	NT2RP2000845	F-NT2RP2000845	0. 78
	NT2RP2000863	F-NT2RP2000863	0.41
35	NT2RP2000880	F-NT2RP2000880	0. 79
	NT2RP2000892	F-NT2RP2000892	0. 94
	NT2RP2000931	F-NT2RP2000931	0. 89
	NT2RP2000932	F-NT2RP2000932	0. 94
40	NT2RP2000938	F-NT2RP2000938	0. 90
40	NT2RP2000943	F-NT2RP2000943	0. 61
	NT2RP2000965	F-NT2RP2000965	0. 80
•	NT2RP2000985	F-NT2RP2000985	0. 94
45	NT2RP2001036	F-NT2RP2001036	0.86
45	NT2RP2001044	F-NT2RP2001044	0. 93
	NT2RP2001056	F-NT2RP2001056	0. 77
	NT2RP2001065		
50	NT2RP2001070	F-NT2RP2001070	0.91
50	NT2RP2001081	F-NT2RP2001081	0.94
	NT2RP2001094	F-NT2RP2001094	0. 93
	NT2RP2001119	F-NT2RP2001119	0.31
55	NT2RP2001127	F-NT2RP2001127	0.41
55	NT2RP2001137	F-NT2RP2001137	0. 93
	NT2RP2001168	F-NT2RP2001168	0.71

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	NT2RP2001173	F-NT2RP2001173	0.94
	NT2RP2001174	F-NT2RP2001174	0.71
	NT2RP2001218	F-NT2RP2001218	0.76
5	NT2RP2001233	F-NT2RP2001233	3
	NT2RP2001245	F-NT2RP2001245	0. 33
	NT2RP2001268	F-NT2RP2001268	0.61
	NT2RP2001327	F-NT2RP2001327	0.86
10			
	NT2RP2001328	F-NT2RP2001328	0.94
	NT2RP2001366	F-NT2RP2001366	0. 52
	NT2RP2001378	F-NT2RP2001378	0.64
	NT2RP2001381	F-NT2RP2001381	0.80
15	NT2RP2001392	F-NT2RP2001392	0.69
	NT2RP2001394	F-NT2RP2001394	0.38
	NT2RP2001397	F-NT2RP2001397	0.79
	NT2RP2001420	F-NT2RP2001420	0.88
20	NT2RP2001427	F-NT2RP2001427	0. 73
	NT2RP2001436	F-NT2RP2001436	0. 32
	NT2RP2001440	F-NT2RP2001440	0.94
	NT2RP2001450	F-NT2RP2001450	0.38
25	NT2RP2001511	F-NT2RP2001511	0.53
	NT2RP2001520	F-NT2RP2001520	0.88
	NT2RP2001536	F-NT2RP2001536	0.64
	NT2RP2001560	F-NT2RP2001560	0.79
30	NT2RP2001576	F-NT2RP2001576	0.62
	NT2RP2001581	F-NT2RP2001581	0.48
	NT2RP2001597	F-NT2RP2001597	0.94
	NT2RP2001601	F-NT2RP2001601	0.33
35	NT2RP2001613	F-NT2RP2001613	0.41
35	NT2RP2001628	F-NT2RP2001628	0.79
	NT2RP2001634	F-NT2RP2001634	0.87
	NT2RP2001660	F-NT2RP2001660	0.86
	NT2RP2001675	F-NT2RP2001675	0.90
40	NT2RP2001721	F-NT2RP2001721	0, 32
	NT2RP2001748	F-NT2RP2001748	0.32
	NT2RP2001839	F-NT2RP2001839	0. 83
	NT2RP2001861	F-NT2RP2001861	0. 33
45	NT2RP2001876	F-NT2RP2001876	0.94
	NT2RP2001883	F-NT2RP2001883	0.90
		F-NT2RP2001898	0.94
	NT2RP2001900	F-NT2RP2001990	0. 93
50	NT2RP2001900	F-NT2RP2001900	0. 93
	NT2RP2001946	F-NT2RP2001946	0.63
	NT2RP2001947	F-NT2RP2001947	0. 75
55	NT2RP2001969	F-NT2RP2001969	0. 94
	NT2RP2001976	F-NT2RP2001976	0. 48
	NT2RP2001985	F-NT2RP2001985	0. 44

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	NT2RP2001991	F-NT2RP2001991	0. 39
	NT2RP2002025	F-NT2RP2002025	0.94
	NT2RP2002033	F-NT2RP2002033	0.40
5	NT2RP2002046	F-NT2RP2002046_	0.56
	NT2RP2002058	F-NT2RP2002058	0.64
	NT2RP2002076	F-NT2RP2002076	0.40
	NT2RP2002078	F-NT2RP2002078	0.81
10	NT2RP2002079	F-NT2RP2002079	0.82
	NT2RP2002099	F-NT2RP2002099	0. 92
	NT2RP2002154	F-NT2RP2002154	0. 44
	NT2RP2002185	F-NT2RP2002185	0.94
15	NT2RP2002193	F-NT2RP2002193	0.48
	NT2RP2002208	F-NT2RP2002208	0.49
	NT2RP2002231	F-NT2RP2002231	
	NT2RP2002235	F-NT2RP2002235	0. 87
20	NT2RP2002252	F-NT2RP2002252	0.51
20	NT2RP2002252	F-NT2RP2002270	0. 78
	NT2RP2002270		
	NT2RP2002292	F-NT2RP2002292 F-NT2RP2002312	0. 94 0. 33
0.5			0. 53
25	NT2RP2002325	F-NT2RP2002325	0. 53
	NT2RP2002333 NT2RP2002373	F-NT2RP2002333	0. 91
		F-NT2RP2002373	
	NT2RP2002385	F-NT2RP2002385	0.94
30	NT2RP2002408	F-NT2RP2002408	0.44
	NT2RP2002426	F-NT2RP2002426	0, 54
	NT2RP2002442	F-NT2RP2002442	0.37
	NT2RP2002464	F-NT2RP2002464	0.79
35	NT2RP2002479	F-NT2RP2002479	0. 92
	NT2RP2002498	F-NT2RP2002498	0.94
	NT2RP2002503	F-NT2RP2002503	0.90
	NT2RP2002520	F-NT2RP2002520	0. 37
40	NT2RP2002537	F-NT2RP2002537	0. 42
	NT2RP2002549	F-NT2RP2002549	0.75
	NT2RP2002595	F-NT2RP2002595	0. 90
	NT2RP2002609	F-NT2RP2002609	0. 33
45	NT2RP2002618	F-NT2RP2002618	0. 52
	NT2RP2002621	F-NT2RP2002621	0. 67
	NT2RP2002672	F-NT2RP2002672	0.43
	NT2RP2002701	F-NT2RP2002701	0.69
50	NT2RP2002706	F-NT2RP2002706	0.51
	NT2RP2002710	F-NT2RP2002710	0.94
	NT2RP2002769	F-NT2RP2002769	0.42
	NT2RP2002800	F-NT2RP2002800	0. 47
E E	NT2RP2002862	F-NT2RP2002862	0.88
55	NT2RP2002880	F-NT2RP2002880	0. 78
	NT2RP2002891	F-NT2RP2002891	0.69

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	NT2RP2002925	F-NT2RP2002925	0.65
	NT2RP2002928	F-NT2RP2002928	0. 47
	NT2RP2002929	F-NT2RP2002929	0.94
5	NT2RP2002939	F-NT2RP2002939	0.94
	NT2RP2002954	F-NT2RP2002954	0.94
	NT2RP2002959	F-NT2RP2002959	0.45
	NT2RP2002979	F-NT2RP2002979	0. 51
10	NT2RP2002980	F-NT2RP2002980	0.38
	NT2RP2002986	F-NT2RP2002986	0. 34
	NT2RP2002993	F-NT2RP2002993	0.94
	NT2RP2003034	F-NT2RP2003034	0. 34
15	NT2RP2003099	F-NT2RP2003099	0. 46
-	NT2RP2003108	F-NT2RP2003108	0. 67
	NT2RP2003117	F-NT2RP2003117,	
	NT2RP2003121	F-NT2RP2003121	0. 94
20	NT2RP2003125	F-NT2RP2003125	0. 94
20	NT2RP2003137	F-NT2RP2003137	0.60
	NT2RP2003157	F-NT2RP2003157	0. 94
	NT2RP2003158	F-NT2RP2003158	0. 75
05	NT2RP2003165	F-NT2RP2003165	0. 90
25	NT2RP2003177	F-NT2RP2003177	0. 85
	NT2RP2003194	F-NT2RP2003194	0. 94
	NT2RP2003228	F-NT2RP2003228	0. 57
	NT2RP2003243	F-NT2RP2003243	0. 94
30	NT2RP2003265	F-NT2RP2003265	0.68
	NT2RP2003272	F-NT2RP2003272	0. 94
	NT2RP2003277	F-NT2RP2003277	0. 73
	NT2RP2003280	F-NT2RP2003280	0.94
35	NT2RP2003286	F-NT2RP2003286	0. 44
	NT2RP2003293	F-NT2RP2003293	0. 42
	NT2RP2003295	F-NT2RP2003295	0.89
	NT2RP2003297	F-NT2RP2003297	0. 94
40	NT2RP2003307	F-NT2RP2003307	0. 79
	NT2RP2003308	F-NT2RP2003308	0. 49
	NT2RP2003329	F-NT2RP2003329	0. 33
	NT2RP2003347	F-NT2RP2003347	0. 79
45	NT2RP2003367	F-NT2RP2003367	0. 94
	NT2RP2003391	F-NT2RP2003391	0. 90
	NT2RP2003393	F-NT2RP2003393	0.94
	NT2RP2003401	F-NT2RP2003401	0. 80
50	NT2RP2003433	F-NT2RP2003433	0. 40
	NT2RP2003445	F-NT2RP2003445	0. 94
	NT2RP2003446	F-NT2RP2003446	0.40
•	NT2RP2003466	F-NT2RP2003466	0. 58
55	NT2RP2003480	F-NT2RP2003480	0.59
	NT2RP2003506	F-NT2RP2003506	0. 94

	NT2RP2003511	F-NT2RP2003511	0: 31
	NT2RP2003513	F-NT2RP2003513	0. 67
	NT2RP2003533	F-NT2RP2003533	0.57
5	NT2RP2003543		0.94
	NT2RP2003564	F-NT2RP2003564	0.88
	NT2RP2003567	F-NT2RP2003567	0.69
	NT2RP2003581	F-NT2RP2003581	0.40
10	NT2RP2003596	F-NT2RP2003596	0.83
	NT2RP2003590	F-NT2RP2003604	0. 87
	NT2RP2003604	F-NT2RP2003629	0.87
	NT2RP2003643	F-NT2RP2003643	0. 94
15	NT2RP2003643	F~NT2RP2003687	0. 31
	NT2RP2003691	F-NT2RP2003691	0.34
	NT2RP2003702	F-NT2RP2003702,	
20	NT2RP2003713 NT2RP2003714	F~NT2RP2003713	0. 53
20	NT2RP2003714	F-NT2RP2003714	0. 61 0. 88
	NT2RP2003737	F-NT2RP2003737 F-NT2RP2003751	0. 36
	NT2RP2003760	F-NT2RP2003760	0. 62
25	NT2RP2003764	F-NT2RP2003764	0. 94
25	NT2RP2003769	F-NT2RP2003769	0. 69
	NT2RP2003777	F-NT2RP2003777	0. 93
	NT2RP2003781	F-NT2RP2003781	0.37
30	NT2RP2003793	F-NT2RP2003793	0.92
30	NT2RP2003825	F-NT2RP2003825	0. 49
	NT2RP2003840	F-NT2RP2003840	0. 88
	NT2RP2003857	F-NT2RP2003857	0.34
35	NT2RP2003952	F-NT2RP2003952	0. 89
33	NT2RP2003976	F-NT2RP2003976	0, 33
	NT2RP2003981	F-NT2RP2003981	0, 64
	NT2RP2003984	F-NT2RP2003984	0.41
40	NT2RP2003986	F-NT2RP2003986	0.32
	NT2RP2004013	F-NT2RP2004013	0.48
	NT2RP2004041	F-NT2RP2004041	0. 78
	NT2RP2004042	F-NT2RP2004042	0.66
45	NT2RP2004066	F-NT2RP2004066	0. 56
40	NT2RP2004081	F-NT2RP2004081	0.94
	NT2RP2004098	F-NT2RP2004098	0.39
	NT2RP2004124	F-NT2RP2004124	0.75
50	NT2RP2004152	F-NT2RP2004152	0.54
	NT2RP2004165	F-NT2RP2004165	0.68
	NT2RP2004187		0.92
	NT2RP2004194		0.90
55	NT2RP2004196	F-NT2RP2004196	0.94
	NT2RP2004226	F-NT2RP2004226	0, 80
	NT2RP2004239	F-NT2RP2004239	0. 63

	NT2RP2004240	F-NT2RP2004240	0. 86
	NT2RP2004242	F-NT2RP2004242	0. 94
_	NT2RP2004245	F-NT2RP2004245	0.92
5	NT2RP2004316	F-NT2RP2004316	_0, 33
	NT2RP2004364	F-NT2RP2004364	0. 75
	NT2RP2004365	F-NT2RP2004365	0.43
	NT2RP2004366	F-NT2RP2004366	0.32
10	NT2RP2004373	F-NT2RP2004373	0.60
•	NT2RP2004389	F-NT2RP2004389	0.94
	NT2RP2004392	F-NT2RP2004392	0. 93
	NT2RP2004399	F-NT2RP2004399	0.36
15	NT2RP2004463	F-NT2RP2004463	0. 43
	NT2RP2004476	F-NT2RP2004476	0. 80
	NT2RP2004538	F-NT2RP2004538	
	NT2RP2004551	F-NT2RP2004551	0. 47
20	NT2RP2004568	F-NT2RP2004568	0. 73
	NT2RP2004594	F-NT2RP2004594	0. 64
	NT2RP2004600	F-NT2RP2004600	0. 53
	NT2RP2004602	F-NT2RP2004602	0. 33
25	NT2RP2004614	F-NT2RP2004614	0. 34
	NT2RP2004655	F-NT2RP2004655	0. 77
	NT2RP2004664	F-NT2RP2004664	0, 94
	NT2RP2004689	F-NT2RP2004689	0. 83
30	NT2RP2004710	F-NT2RP2004710	0. 94
••	NT2RP2004743	F-NT2RP2004743	0. 32
	NT2RP2004768		0. 53
	NT2RP2004791	F-NT2RP2004791	0. 64
35	NT2RP2004799	F-NT2RP2004799	0. 42
35	NT2RP2004802	F-NT2RP2004802	0. 88
	NT2RP2004816	F-NT2RP2004816	0. 78
	NT2RP2004841	F-NT2RP2004841	0. 61
40	NT2RP2004861	F-NT2RP2004861	0. 32
40	NT2RP2004897	F-NT2RP2004897	0. 67
	NT2RP2004933	F-NT2RP2004933	0. 94
	NT2RP2004936	F-NT2RP2004936	0. 53
	NT2RP2004959	F-NT2RP2004959	0. 78
45	NT2RP2004978	F-NT2RP2004978	0. 86
	NT2RP2004985	F-NT2RP2004985	0. 42
	NT2RP2004999	F-NT2RP2004999	0. 55
	NT2RP2005000	F-NT2RP2005000	0. 40
50	NT2RP2005001	F-NT2RP2005001	0. 94
	NT2RP2005003	F-NT2RP2005003	0. 44
	NT2RP2005012	F-NT2RP2005012	0. 73
	NT2RP2005018	F-NT2RP2005018	0. 43
55	NT2RP2005022	F-NT2RP2005022	0. 34
	NT2RP2005037	F-NT2RP2005037	0. 94
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	NT2RP2005038	F-NT2RP2005038	0.56
	NT2RP2005116	F-NT2RP2005116	0. 94
	NT2RP2005126	F-NT2RP2005126	0. 69
5	NT2RP2005139	F-NT2RP2005139	0.94
	NT2RP2005140	F-NT2RP2005140	0. 32
	NT2RP2005144	F-NT2RP2005144	0.39
	NT2RP2005147	F-NT2RP2005147	0. 32
10	NT2RP2005159	F-NT2RP2005159	0. 52
	NT2RP2005162	F-NT2RP2005162	0. 86
	NT2RP2005168	F-NT2RP2005168	0. 53
	NT2RP2005204	F-NT2RP2005204	0. 94
15	NT2RP2005227	F-NT2RP2005227	0. 53
10	NT2RP2005239	F-NT2RP2005239	0. 91
	NT2RP2005270	F-NT2RP2005270,	
	NT2RP2005276	F-NT2RP2005276	0. 41
	NT2RP2005287	F-NT2RP2005287	0. 44
20	NT2RP2005288	F-NT2RP2005288	0. 82
	NT2RP2005315	F-NT2RP2005315	0. 79
	NT2RP2005358	F-NT2RP2005358	0. 94
	NT2RP2005393	F-NT2RP2005393	0. 41
25	NT2RP2005436	F-NT2RP2005436	0. 45
	NT2RP2005441	F-NT2RP2005441	0. 88
	NT2RP2005453	F-NT2RP2005453	0. 59
	NT2RP2005457	F-NT2RP2005457	0. 56
30	NT2RP2005464	F-NT2RP2005464	0. 72
	NT2RP2005465	F-NT2RP2005465	0.40
	NT2RP2005472	F-NT2RP2005472	0. 42
	NT2RP2005490	F-NT2RP2005490	0, 52
35	NT2RP2005495	F-NT2RP2005495	0. 94
	NT2RP2005496	F-NT2RP2005496	0. 52
	NT2RP2005498	F-NT2RP2005498	0. 74
	NT2RP2005509	F-NT2RP2005509	0. 55
40	NT2RP2005520	F-NT2RP2005520	0. 54
	NT2RP2005525	F-NT2RP2005525	0. 41
	NT2RP2005539	F-NT2RP2005539	0. 94
	NT2RP2005540	F-NT2RP2005540	0. 32
45	NT2RP2005549	F-NT2RP2005549	0. 79
	NT2RP2005555	F-NT2RP2005555	0. 63
	NT2RP2005557	F-NT2RP2005557	0, 79
	NT2RP2005600	F-NT2RP2005600	0. 76
50	NT2RP2005605	F-NT2RP2005605	0.56
	NT2RP2005620	F-NT2RP2005620	0.66
	NT2RP2005622	F-NT2RP2005622	0.37
	NT2RP2005635	F-NT2RP2005635	0. 87
55	NT2RP2005637	F-NT2RP2005637	0.94
	NT2RP2005640	F-NT2RP2005640	0. 87

	NT2RP2005654	F-NT2RP2005654	0.94
	NT2RP2005669	F-NT2RP2005669	0. 94
	NT2RP2005675	F-NT2RP2005675	0. 72
5	NT2RP2005683	======	0.72
3		F-NT2RP2005683	•
	NT2RP2005690	F-NT2RP2005690	0.54
	NT2RP2005712	F-NT2RP2005712	0.93
	NT2RP2005722	F-NT2RP2005722	0.50
10	NT2RP2005723	F-NT2RP2005723	0. 45
	NT2RP2005732	F-NT2RP2005732	0. 54
	NT2RP2005748	F-NT2RP2005748	0.76
	NT2RP2005752	F-NT2RP2005752	0. 47
15	NT2RP2005763	F-NT2RP2005763	0. 87
	NT2RP2005767	F-NT2RP2005767	0. 58
	NT2RP2005773	F-NT2RP2005773,	
	NT2RP2005775	F-NT2RP2005775	0.34
20	NT2RP2005781	F-NT2RP2005781	0.81
	NT2RP2005784	F-NT2RP2005784	0.94
	NT2RP2005804	F-NT2RP2005804	0.94
	NT2RP2005812	F-NT2RP2005812	0.90
25	NT2RP2005835	F-NT2RP2005835	0.94
	NT2RP2005853	F-NT2RP2005853	0.77
	NT2RP2005859	F-NT2RP2005859	0.68
	NT2RP2005868	F-NT2RP2005868	0.76
30	NT2RP2005886	F-NT2RP2005886	0.62
•••	NT2RP2005890	F-NT2RP2005890	0.79
	NT2RP2005901	F-NT2RP2005901	0.94
	NT2RP2005933	F-NT2RP2005933	0. 55
35	NT2RP2006023	F-NT2RP2006023	0. 47
39	NT2RP2006038	F-NT2RP2006038	0.73
	NT2RP2006043	F-NT2RP2006043	0.50
	NT2RP2006052	F-NT2RP2006052	0.74
	NT2RP2006069	F-NT2RP2006069	0.85
40	NT2RP2006071	F-NT2RP2006071	0. 94
	NT2RP2006100	F-NT2RP2006100	0. 42
	NT2RP2006106	F-NT2RP2006106	0. 94
	NT2RP2006141	F-NT2RP2006141	0. 92
45	NT2RP2006186	F-NT2RP2006186	0.94
	NT2RP2006196	F-NT2RP2006196	0.32
	NT2RP2006200	F-NT2RP2006200	0. 70
	NT2RP2006219	F-NT2RP2006219	0.94
50	NT2RP2006237	F-NT2RP2006237	0.59
	NT2RP2006238	F-NT2RP2006238	0. 47
	NT2RP2006275	F-NT2RP2006275	0.89
	NT2RP2006312	F-NT2RP2006312	0.75
55	NT2RP2006333	F-NT2RP2006333	0. 73
	NT2RP2006334	F-NT2RP2006334	0.72

	NT2RP2006365	F-NT2RP2006365	0. 48
	NT2RP2006436	F-NT2RP2006436	0. 75
	NT2RP2006441	F-NT2RP2006441	0. 73
5	NT2RP2006456	F-NT2RP2006456	0. 60
	NT2RP2006464	F-NT2RP2006464	0.51
	NT2RP2006472		
		F-NT2RP2006472	0. 76
10	NT2RP2006565	F-NT2RP2006565	0. 34
70	NT2RP2006571	F-NT2RP2006571	0. 94
	NT2RP2006573	F-NT2RP2006573	0. 35
	NT2RP3000002	F-NT2RP3000002	0.40
45	NT2RP3000031	F-NT2RP3000031	0. 49
15	NT2RP3000046	F-NT2RP3000046	0. 90
	NT2RP3000047	F-NT2RP3000047	0. 69
	NT2RP3000050	F-NT2RP3000050	
	NT2RP3000068	F-NT2RP3000068	0. 94
20	NT2RP3000072	F-NT2RP3000072	0.34
	NT2RP3000080	F-NT2RP3000080	0. 32
	NT2RP3000085	F-NT2RP3000085	0. 92
•	NT2RP3000092	F-NT2RP3000092	0. 40
25	NT2RP3000134	F-NT2RP3000134	0. 42
	NT2RP3000142	F-NT2RP3000142	0. 58
	NT2RP3000149	F-NT2RP3000149	0. 83
	NT2RP3000197	F-NT2RP3000197	0. 59
30	NT2RP3000207	F-NT2RP3000207	0. 48
	NT2RP3000220	F-NT2RP3000220	0. 75
	NT2RP3000251	F-NT2RP3000251	0. 82
	NT2RP3000252	F-NT2RP3000252	0. 88
35	NT2RP3000255	F-NT2RP3000255	0. 94
	NT2RP3000267	F-NT2RP3000267	0. 92
	NT2RP3000299	F-NT2RP3000299	0. 94
	NT2RP3000312	F-NT2RP3000312	0. 57
40	NT2RP3000320	F-NT2RP3000320	0. 94
	NT2RP3000324	F-NT2RP3000324	0. 32
	NT2RP3000333	F-NT2RP3000333	0. 94
	NT2RP3000348	F-NT2RP3000348	0. 32
45	NT2RP3000350	F-NT2RP3000350	0. 67
45	NT2RP3000359	F-NT2RP3000359	0.94
	NT2RP3000361	F-NT2RP3000361	0. 94
	NT2RP3000366	F-NT2RP3000366	0. 94
	NT2RP3000393	F-NT2RP3000393	0. 53
50	NT2RP3000397	F-NT2RP3000397	0.89
	NT2RP3000403	F-NT2RP3000403	0.87
	NT2RP3000439	F-NT2RP3000439	0. 77
•	NT2RP3000456	F-NT2RP3000456	0.88
55	NT2RP3000484	F-NT2RP3000484	0. 59
	NT2RP3000487	F-NT2RP3000487	0. 32

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	NT2RP3000526	F-NT2RP3000526	0. 32
	NT2RP3000527	F-NT2RP3000527	0. 69
	NT2RP3000531	F-NT2RP3000531	0. 56
5	NT2RP3000562	F-NT2RP3000562	0. 44
	NT2RP3000590	F-NT2RP3000590	0. 88
	NT2RP3000592	F-NT2RP3000592	0. 92
	NT2RP3000596	F-NT2RP3000596	0. 52
10	NT2RP3000599		0. 69
70		F-NT2RP3000599	
	NT2RP3000603	F-NT2RP3000603	0. 94
	NT2RP3000605	F-NT2RP3000605	0. 92
	NT2RP3000624	F-NT2RP3000624	0. 71
15	NT2RP3000632	F-NT2RP3000632	0. 43
	NT2RP3000644	F-NT2RP3000644	0. 94
	NT2RP3000661	F-NT2RP3000661,	0.60
	NT2RP3000665	F-NT2RP3000665	0. 46
20	NT2RP3000690	F-NT2RP3000690	0. 79
	NT2RP3000739	F-NT2RP3000739	0.94
	NT2RP3000753	F-NT2RP3000753	0. 42
	NT2RP3000759	F-NT2RP3000759	0. 94
25	NT2RP3000825	F-NT2RP3000825	0.73
	NT2RP3000826	F-NT2RP3000826	0. 94
	NT2RP3000836	F-NT2RP3000836	0. 39
	NT2RP3000841	F-NT2RP3000841	0. 55
20	NT2RP3000845	F-NT2RP3000845	0, 38
30	NT2RP3000847		0. 67
	NT2RP3000850		0. 59
	NT2RP3000852		0, 39
	NT2RP3000859	F-NT2RP3000859	0. 83
35	NT2RP3000868	F-NT2RP3000868	0. 49
	NT2RP3000869	F-NT2RP3000869	0. 94
	NT2RP3000901		0. 34
	NT2RP3000917	F-NT2RP3000917	0. 73
40	NT2RP3000919	F-NT2RP3000919	0. 76
	NT2RP3000968	F-NT2RP3000968	0. 94
	NT2RP3000980	F-NT2RP3000980	0. 61
	NT2RP3000994	F-NT2RP3000994	0. 45
45	NT2RP3001004	F-NT2RP3001004	0. 45
	NT2RP3001004	F-NT2RP3001004	0. 38
	NT2RP3001057	F-NT2RP3001057	_
	NT2RP3001057	F-NT2RP3001057	0. 84 0. 57
50	NT2RP3001081		
		F-NT2RP3001084	0.94
	NT2RP3001096	F-NT2RP3001096	0. 94
	NT2RP3001107	F-NT2RP3001107	0. 88
55	NT2RP3001109	F-NT2RP3001109	0.67
55	NT2RP3001111	F-NT2RP3001111	0.87
	NT2RP3001113	F-NT2RP3001113	0. 40

	NT2RP3001116	F-NT2RP3001116	0. 73
	NT2RP3001119	F-NT2RP3001119	0. 62
	NT2RP3001120	F-NT2RP3001120	0. 59
5	NT2RP3001126	F-NT2RP3001126	0. 94
	NT2RP3001133	F-NT2RP3001133	0, 89
	NT2RP3001140	F-NT2RP3001140	0. 59
	NT2RP3001147	F-NT2RP3001147	0. 94
10	NT2RP3001150	F-NT2RP3001150	0. 92
•	NT2RP3001155	F-NT2RP3001155	0. 94
	NT2RP3001176	F-NT2RP3001176	0. 82
	NT2RP3001214	F-NT2RP3001214	0. 41
15	NT2RP3001216	F-NT2RP3001216	0. 32
	NT2RP3001221	F-NT2RP3001221	0. 46
	NT2RP3001236	F-NT2RP3001236,	
	NT2RP3001239	F-NT2RP3001239	0. 73
20	NT2RP3001253	F-NT2RP3001253	0. 59
20	NT2RP3001260	F-NT2RP3001260	0. 94
	NT2RP3001268	F-NT2RP3001268	0. 75
	NT2RP3001278	F-NT2RP3001272	0. 75
0.5	NT2RP3001274	F-NT2RP3001274	0. 94
25	NT2RP3001297	F-NT2RP3001274	0. 94
	NT2RP3001307	F-NT2RP3001307	0. 87
	NT2RP3001307	F-NT2RP3001307	0. 87
	NT2RP3001328	F-NT2RP3001328	0. 71
30	NT2RP3001355	F-NT2RP3001355	0. 40
	NT2RP3001356	F-NT2RP3001356	0. 40
	NT2RP3001384	F-NT2RP3001384	0. 91
	NT2RP3001392	F-NT2RP3001392	0. 42
35	NT2RP3001392	F-NT2RP3001392	0. 42
	NT2RP3001398	F-NT2RP3001398	0. 94
	NT2RP3001399	F-NT2RP3001399	0. 94
	NT2RP3001407	F-NT2RP3001407	0. 54
40	NT2RP3001407	F-NT2RP3001407	0. 74
•	NT2RP3001426	F-NT2RP3001426	0. 74
	NT2RP3001427	F-NT2RP3001427	0. 92
	NT2RP3001427	F-NT2RP3001427	0. 40
45	NT2RP3001447	F-NT2RP3001428	0. 57
		F-NT2RP3001447	
	NT2RP3001449		0. 62
	NT2RP3001453	F-NT2RP3001453	0. 94
50	NT2RP3001457	F-NT2RP3001457	0.89
	NT2RP3001472	F-NT2RP3001472	0. 82
	NT2RP3001490	F-NT2RP3001490	0. 53
	NT2RP3001495	F-NT2RP3001495	0. 94
55	NT2RP3001497	F-NT2RP3001497	0.40
	NT2RP3001529	F-NT2RP3001529	0.94
	NT2RP3001538	F-NT2RP3001538	0. 93
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	NT-2RP3001554	F-NT2RP3001554	0. 91
	NT2RP3001587	F-NT2RP3001587	0. 94
	NT2RP3001608	F-NT2RP3001608	0. 94
5	NT2RP3001621		
	NT2RP3001629	F-NT2RP3001621_ F-NT2RP3001629	0. 78 0. 42
	NT2RP3001642		
		F-NT2RP3001642	0.94
10	NT2RP3001646	F-NT2RP3001646	0.94
	NT2RP3001671	F-NT2RP3001671	0.77
	NT2RP3001672	F-NT2RP3001672	0.88
	NT2RP3001676	F-NT2RP3001676	0. 88
	NT2RP3001678	F-NT2RP3001678	0.44
15	NT2RP3001679	F-NT2RP3001679	0. 44
	NT2RP3001688	F-NT2RP3001688	0. 67
	NT2RP3001690	F-NT2RP3001690	0.58
	NT2RP3001698	F-NT2RP3001698	0.86
20	NT2RP3001708	F-NT2RP3001708	0.38
	NT2RP3001712	F-NT2RP3001712	0.75
	NT2RP3001716	F-NT2RP3001716	0.77
	NT2RP3001724	F-NT2RP3001724	0.65
25	NT2RP3001727	F-NT2RP3001727	0.84
	NT2RP3001730	F-NT2RP3001730	0. 77
	NT2RP3001739	F-NT2RP3001739	0. 78
	NT2RP3001753	F-NT2RP3001753	0. 79
30	NT2RP3001764	F-NT2RP3001764	0. 33
	NT2RP3001777	F-NT2RP3001777	0. 87
	NT2RP3001792	F-NT2RP3001792	0.86
	NT2RP3001799	F-NT2RP3001799	0. 79
35	NT2RP3001819	F-NT2RP3001819	0. 31
	NT2RP3001844	F-NT2RP3001844	0. 34
	NT2RP3001854	F-NT2RP3001854	0. 94
	NT2RP3001855	F-NT2RP3001855	0.88
40	NT2RP3001857	F-NT2RP3001857	0. 41
	NT2RP3001896	F-NT2RP3001896	0. 94
	NT2RP3001915	F-NT2RP3001915	0. 42
	NT2RP3001929	F-NT2RP3001929	0. 49
45	NT2RP3001931	F-NT2RP3001931	0. 94
40	NT2RP3001938	F-NT2RP3001938	0. 32
	NT2RP3001943	F-NT2RP3001943	0. 70
	NT2RP3001944	F-NT2RP3001944	0. 94
50	NT2RP3001969	F-NT2RP3001969	0. 54
50	NT2RP3002007	F-NT2RP3002007	0. 56
	NT2RP3002014	F-NT2RP3002014	0. 94
	NT2RP3002033	F-NT2RP3002033	0. 94
	NT2RP3002045	F-NT2RP3002045	0.91
55	NT2RP3002054	F-NT2RP3002054	0. 94
	NT2RP3002062	F-NT2RP3002062	0. 62

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	NT2RP3002063	F-NT2RP3002063	0.34
	NT2RP3002081	F-NT2RP3002081	0.57
	NT2RP3002097	F-NT2RP3002097	0.72
5	NT2RP3002102	F-NT2RP3002102	0.90
	NT2RP3002108	F-NT2RP3002108	0.74
	NT2RP3002142	F-NT2RP3002142	0.61
	NT2RP3002146	F-NT2RP3002146	0.91
10	NT2RP3002147	F-NT2RP3002147	0.37
	NT2RP3002151	F-NT2RP3002151	0.90
	NT2RP3002151	F-NT2RP3002163	0. 76
			0. 70
45	NT2RP3002165	F-NT2RP3002165	
15	NT2RP3002181	F-NT2RP3002181	0. 78
	NT2RP3002244	F-NT2RP3002244	0.81
	NT2RP3002248	F-NT2RP3002248,	
	NT2RP3002255	F-NT2RP3002255	0. 53
20	NT2RP3002273	F-NT2RP3002273	0. 90
	NT2RP3002276	F-NT2RP3002276	0. 39
	NT2RP3002303	F-NT2RP3002303	0. 40
	NT2RP3002304	F-NT2RP3002304	0. 44
25	NT2RP3002330	F-NT2RP3002330	0.94
	NT2RP3002343	F-NT2RP3002343	0.94
	NT2RP3002351	F-NT2RP3002351	0.94
	NT2RP3002377	F-NT2RP3002377	0.52
30	NT2RP3002399	F-NT2RP3002399	0. 91
	NT2RP3002402	F-NT2RP3002402	0. 33
	NT2RP3002455	F-NT2RP3002455	0. 42
	NT2RP3002484	F-NT2RP3002484	0. 52
35	NT2RP3002501	F-NT2RP3002501	0. 57
	NT2RP3002512	F-NT2RP3002512	0. 63
	NT2RP3002529	F-NT2RP3002529	0.74
	NT2RP3002545	F-NT2RP3002545	0. 94
40	NT2RP3002549	F-NT2RP3002549	0. 94
40	NT2RP3002566	F-NT2RP3002566	0.74
	NT2RP3002587	F-NT2RP3002587	0.94
	NT2RP3002590	F-NT2RP3002590	0. 37
	NT2RP3002602	F-NT2RP3002602	0.39
45	NT2RP3002603	F-NT2RP3002603	0.46
	NT2RP3002628	F-NT2RP3002628	0.31
	NT2RP3002631	F-NT2RP3002631	0. 79
	NT2RP3002650	F-NT2RP3002650	0. 94
50	NT2RP3002659	F-NT2RP3002659	0. 93
	NT2RP3002660	F-NT2RP3002660	0. 46
	NT2RP3002663	F-NT2RP3002663	0.79
	NT2RP30026671	F-NT2RP3002663	0.79
55	NT2RP3002671	F-NT2RP3002682	0. 52
	NT2RP3002682	F-NT2RP3002688	0. 52
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	NT2RP3002701	F-NT2RP3002701	0. 62
	NT2RP3002763	F-NT2RP3002763	0. 83
	NT2RP3002785	F-NT2RP3002785	0. 33
5	NT2RP3002818	F-NT2RP3002818	0. 91
	NT2RP3002861	F-NT2RP3002861	0.31
	NT2RP3002869	F-NT2RP3002869	0. 47
	NT2RP3002876	F-NT2RP3002876	0. 58
10	NT2RP3002877	F-NT2RP3002877	0. 41
	NT2RP3002909	F-NT2RP3002909	0. 52
	NT2RP3002911	F-NT2RP3002911	0. 73
	NT2RP3002948	F-NT2RP3002948	0. 60
15	NT2RP3002953	F-NT2RP3002953	0. 37
,,	NT2RP3002969	F-NT2RP3002969	0. 37
	NT2RP3002972	F-NT2RP3002972,	
	NT2RP3002985	F-NT2RP3002985	0. 55
20	NT2RP3002988	F-NT2RP3002988	0. 82
20	NT2RP3003008	F-NT2RP3003008	0. 94
	NT2RP3003061	F-NT2RP3003061	0. 93
	NT2RP3003061	F-NT2RP3003068	0. 45
05	NT2RP3003071	F-NT2RP3003071	0. 76
25	NT2RP3003078	F-NT2RP3003078	0. 40
	NT2RP3003101	F-NT2RP3003101	0. 40
	NT2RP3003139	F-NT2RP3003139	0.41
	NT2RP3003145	F-NT2RP3003145	0. 43
30	NT2RP3003150	F-NT2RP3003150	0.94
	NT2RP3003157	F-NT2RP3003157	0. 94
	NT2RP3003185	F-NT2RP3003185	0. 87
	NT2RP3003193	F-NT2RP3003193	0. 94
35	NT2RP3003197	F-NT2RP3003197	0. 74
	NT2RP3003203	F-NT2RP3003203	0. 94
	NT2RP3003204	F-NT2RP3003204	0. 74
	NT2RP3003210	F-NT2RP3003210	0. 45
40	NT2RP3003212	F-NT2RP3003212	0. 51
	NT2RP3003230	F-NT2RP3003230	0. 94
	NT2RP3003242	F-NT2RP3003242	0. 92
	NT2RP3003251	F-NT2RP3003251	0. 93
45	NT2RP3003278	F-NT2RP3003278	0. 83
	NT2RP3003282	F-NT2RP3003282	0. 67
	NT2RP3003290	F-NT2RP3003290	0. 62
	NT2RP3003301	F-NT2RP3003301	0. 49
50	NT2RP3003302	F-NT2RP3003302	0.32
	NT2RP3003311	F-NT2RP3003311	0. 34
	NT2RP3003313	F-NT2RP3003313	0.88
	NT2RP3003327	F-NT2RP3003327	0. 88
55	NT2RP3003330	F-NT2RP3003330	0. 94
	NT2RP3003344	F-NT2RP3003344	0. 67

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	NT2RP3003353	F-NT2RP3003353	0. 65
	NT2RP3003377	F-NT2RP3003377	0.87
	NT2RP3003385	F-NT2RP3003385	0. 94
5	NT2RP3003409	F-NT2RP3003409	0.41
	NT2RP3003411	F-NT2RP3003411	0. 47
	NT2RP3003427	F-NT2RP3003427	0. 33
	NT2RP3003490	F-NT2RP3003490	0. 56
10	NT2RP3003491	F-NT2RP3003491	0.37
	NT2RP3003543		0. 92
			0. 64
	NT2RP3003552		
	NT2RP3003555		0. 94
15		F-NT2RP3003564	0. 86
		F-NT2RP3003589	0. 88
		F-NT2RP3003621,	
	NT2RP3003656	F-NT2RP3003656	0. 77
20	NT2RP3003659	F-NT2RP3003659	0. 32
	NT2RP3003672	F-NT2RP3003672	0.49
	NT2RP3003680	F-NT2RP3003680	0. 94
	NT2RP3003686	F-NT2RP3003686	0. 78
25	NT2RP3003701	F-NT2RP3003701	0.36
	NT2RP3003716	F-NT2RP3003716	0. 34
	NT2RP3003726	F-NT2RP3003726	0.72
	NT2RP3003795	F-NT2RP3003795	0. 34
30	NT2RP3003805	F-NT2RP3003805	0. 74
	NT2RP3003809	F-NT2RP3003809	0. 73
	NT2RP3003825	F-NT2RP3003825	0. 94
	NT2RP3003831	F-NT2RP3003831	0. 52
35	NT2RP3003833	F-NT2RP3003833	0.31
30	NT2RP3003846	F-NT2RP3003846	0. 54
	NT2RP3003870	F-NT2RP3003870	0. 91
	NT2RP3003876	F-NT2RP3003876	0. 42
	NT2RP3003914	F-NT2RP3003914	0.94
40	NT2RP3003918	F-NT2RP3003918	0. 90
	NT2RP3003932	F-NT2RP3003932	0. 64
	NT2RP3003992	F-NT2RP3003992	0.94
		F-NT2RP3004013	0.60
45	NT2RP3004041		0. 52
	NT2RP3004051	F-NT2RP3004051	0. 75
	NT2RP3004078	F-NT2RP3004078	0. 73
	NT2RP3004093	F-NT2RP3004093	0. 64
50	NT2RP3004125	F-NT2RP3004125	0. 94
	NT2RP3004155	F-NT2RP3004155	0. 89
	NT2RP3004189		0. 46
	NT2RP3004206		0. 59
55	NT2RP3004206		0. 39
	NT2RP3004209	F-NT2RP3004209	0. 54
	M12RF3004203	1 -111214-3004209	0, 54
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		01401174	
	NT2RP3004242	F-NT2RP3004242	0.61
	NT2RP3004246	F-NT2RP3004246	0.91
	NT2RP3004253	F-NT2RP3004253	0.41
5	NT2RP3004258	F-NT2RP3004258	0.41
	NT2RP3004262	F-NT2RP3004262	0.56
	NT2RP3004282	F-NT2RP3004282	0.90
	NT2RP3004332	F-NT2RP3004332	0. 53
10	NT2RP3004341	F-NT2RP3004341	0.40
	NT2RP3004348	F-NT2RP3004348	0.94
	NT2RP3004378	F-NT2RP3004378	0.40
	NT2RP3004424	F-NT2RP3004424	0. 75
15	NT2RP3004424	F-NT2RP3004428	0. 75
75	NT2RP3004451	F-NT2RP3004451	0.85
	NT2RP3004454	F-NT2RP3004454,	
		•	0. 93
•	NT2RP3004466	F-NT2RP3004466 F-NT2RP3004472	0. 80
20	NT2RP3004472	F-NT2RP3004472	0. 40
	NT2RP3004480		0. 77
	NT2RP3004490	F-NT2RP3004490 F-NT2RP3004498	
	NT2RP3004498 NT2RP3004504		0. 91 0. 88
25		F-NT2RP3004504	
	NT2RP3004507	F-NT2RP3004507	0. 86 0. 32
	NT2RP3004534	F-NT2RP3004534	****
	NT2RP3004539	F-NT2RP3004539	0.76
30	NT2RP3004544	F-NT2RP3004544	0.94
	NT2RP3004566	F-NT2RP3004566	0.94
	NT2RP3004569		0.69
	NT2RP3004572		0.65
35	NT2RP3004578	F-NT2RP3004578	0. 52
	NT2RP3004594	F-NT2RP3004594	0.39
	NT2RP3004617	F-NT2RP3004617	0. 94
	NT2RP3004618	F-NT2RP3004618	0.41
40	NT2RP3004669	F-NT2RP3004669	0. 82
	NT2RP3004670	F-NT2RP3004670	0. 32
	NT2RP4000008	F-NT2RP4000008	0. 94
	NT2RP4000051	F-NT2RP4000051	0. 94
45	NT2RP4000109	F-NT2RP4000109	0. 84
	NT2RP4000111	F-NT2RP4000111	0.49
	NT2RP4000129	F-NT2RP4000129	0, 94
	NT2RP4000147	F-NT2RP4000147	0. 92
50	NT2RP4000150	F-NT2RP4000150	0. 41
-	NT2RP4000151	F-NT2RP4000151	0. 33
	NT2RP4000159	F-NT2RP4000159	0.75
	NT2RP4000185	F-NT2RP4000185	0. 94
55	NT2RP4000210	F-NT2RP4000210	0. 83
	NT2RP4000212	F-NT2RP4000212	0. 94
	NT2RP4000243	F-NT2RP4000243	0. 94

	NT2RP4000246-	F-NT2RP4000246	0. 59
	NT2RP4000259	F-NT2RP4000259	0. 94
	NT2RP4000263	F-NT2RP4000263	0. 62
5	NT2RP4000290	F-NT2RP4000290_	0. 92
	NT2RP4000312	F-NT2RP4000312	0.39
·	NT2RP4000323	F-NT2RP4000323	0.73
	NT2RP4000355	F-NT2RP4000355	0.94
10	NT2RP4000360	F-NT2RP4000360	0. 55
	NT2RP4000367	F-NT2RP4000367	0.60
	NT2RP4000370	F-NT2RP4000370	0. 93
	NT2RP4000376	F-NT2RP4000376	0.88
15	NT2RP4000381	F-NT2RP4000381	0. 39
	NT2RP4000398	F-NT2RP4000398	0. 64
	NT2RP4000415	F-NT2RP4000415,	
	NT2RP4000417	F-NT2RP4000417	0. 33
20	NT2RP4000448	F-NT2RP4000448	0. 33
	NT2RP4000449	F-NT2RP4000449	0. 33
	NT2RP4000455	F-NT2RP4000455	0. 37
	NT2RP4000457	F-NT2RP4000457	0.36
25	NT2RP4000480	F-NT2RP4000480	0.88
	NT2RP4000481	F-NT2RP4000481	0. 51 0. 64
	NT2RP4000498 NT2RP4000500	F-NT2RP4000498 F-NT2RP4000500	0. 67
	NT2RP4000500	F-NT2RP4000518	0. 34
30	NT2RP4000524	F-NT2RP4000524	0. 94
	NT2RP4000528	F-NT2RP4000528	0. 65
	NT2RP4000541	F-NT2RP4000541	0. 44
25	NT2RP4000556	F-NT2RP4000556	0. 69
35	NT2RP4000560	F-NT2RP4000560	0. 67
	NT2RP4000588	F-NT2RP4000588	0. 9 4
	NT2RP4000614	F-NT2RP4000614	0. 94
40	NT2RP4000638	F-NT2RP4000638	0. 74
70	NT2RP4000648	F-NT2RP4000648	0. 40
	NT2RP4000657	F-NT2RP4000657	0. 94
	NT2RP4000704	F-NT2RP4000704	0. 69
45	NT2RP4000713	F-NT2RP4000713	0. 93
	NT2RP4000724	F-NT2RP4000724	0. 32
	NT2RP4000728	F-NT2RP4000728	0. 34
	NT2RP4000737	F-NT2RP4000737	0. 48
50	NT2RP4000739	F-NT2RP4000739	0. 40
	NT2RP4000781	F-NT2RP4000781	0. 94
	NT2RP4000787	F-NT2RP4000787	0. 94
	NT2RP4000817 NT2RP4000833	F-NT2RP4000817	0. 60 0. 35
55	NT2RP4000833	F-NT2RP4000833 F-NT2RP4000837	0. 80
	NT2RP4000837	F-NT2RP4000839	0. 33
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	NTODDAGGGG	E NT0004000EE	0.74
	NT2RP4000855	F-NT2RP4000855	0. 74
	NT2RP4000865	F-NT2RP4000865	0. 77
_	NT2RP4000878	F-NT2RP4000878	0. 79
5	NT2RP4000879	F-NT2RP4000879	•
	NT2RP4000907	F-NT2RP4000907	0. 44
	NT2RP4000918	F-NT2RP4000918	0. 88
	NT2RP4000925	F-NT2RP4000925	0. 89
10	NT2RP4000927	F-NT2RP4000927	0. 81
	NT2RP4000928	F-NT2RP4000928	0. 39
	NT2RP4000929	F-NT2RP4000929	0. 34
	NT2RP4000955	F-NT2RP4000955	0. 77
15	NT2RP4000973	F-NT2RP4000973	0. 36
	NT2RP4000975	F-NT2RP4000975	0. 87
	NT2RP4000979	F-NT2RP4000979,	0. 50
	NT2RP4000984	F-NT2RP4000984	0. 66
20	NT2RP4000989	F-NT2RP4000989	0. 94
	NT2RP4000997	F-NT2RP4000997	0. 68
	NT2RP4001004	F-NT2RP4001004	0.40
	NT2RP4001006	F-NT2RP4001006	0. 91
25	NT2RP4001010	F-NT2RP4001010	0. 64
	NT2RP4001029	F-NT2RP4001029	0. 77
	NT2RP4001041	F-NT2RP4001041	0. 79
•	NT2RP4001057	F-NT2RP4001057	0, 65
30	NT2RP4001064	F-NT2RP4001064	0. 94
	NT2RP4001078	F-NT2RP4001078	0. 39
	NT2RP4001079	F-NT2RP4001079	0. 94
	NT2RP4001080	F-NT2RP4001080	0. 58
35	NT2RP4001086	F-NT2RP4001086	0. 78
	NT2RP4001095	F-NT2RP4001095	0. 80
	NT2RP4001100	F-NT2RP4001100	0. 79
	NT2RP4001117	F-NT2RP4001117	0. 60
40	NT2RP4001122	F-NT2RP4001122	0.82
70	NT2RP4001126	F-NT2RP4001126	0. 55
	NT2RP4001138	F-NT2RP4001138	0. 94
	NT2RP4001143	F-NT2RP4001143	0. 41
45	NT2RP4001148	F-NT2RP4001148	0. 37
45	NT2RP4001149	F-NT2RP4001149	0. 50
	NT2RP4001150	F-NT2RP4001150	0. 94
	NT2RP4001159	F-NT2RP4001159	0. 78
50	NT2RP4001174	F-NT2RP4001174	0.94
50	NT2RP4001206	F-NT2RP4001206	0. 80
	NT2RP4001207	F-NT2RP4001207	0. 38
	NT2RP4001210	F-NT2RP4001210	0. 94
	NT2RP4001213	F-NT2RP4001213	0. 46
55	NT2RP4001219	F-NT2RP4001219	0. 38
	NT2RP4001228	F-NT2RP4001228	0. 61
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		0.4011712	
	NT2RP4001235	F-NT2RP4001235	0.88
	NT2RP4001256	F-NT2RP4001256	0.60
	NT2RP4001260	F-NT2RP4001260	0. 94
5	NT2RP4001274	F-NT2RP4001274	0. 82
	NT2RP4001276	F-NT2RP4001276	0. 68
	NT2RP4001313	F-NT2RP4001313	0.77
	NT2RP4001315	F-NT2RP4001315	0. 87
10	NT2RP4001336	F-NT2RP4001336	0. 48
	NT2RP4001339	F-NT2RP4001339	0. 41
	NT2RP4001343	F-NT2RP4001343	0. 63
	NT2RP4001345	F-NT2RP4001345	0. 89
15	NT2RP4001351	F-NT2RP4001351	0. 87
	NT2RP4001353	F-NT2RP4001353	0.72
	NT2RP4001372	F-NT2RP4001372	
	NT2RP4001373	F-NT2RP4001373	0. 60
20	NT2RP4001375	F-NT2RP4001375	0.43
20	NT2RP4001379	F-NT2RP4001379	0. 77
	NT2RP4001389	F-NT2RP4001389	0. 77
	NT2RP4001414	F-NT2RP4001414	0. 90
25	NT2RP4001414	F-NT2RP4001433	0. 39
25	NT2RP4001442	F-NT2RP4001442	0. 59
	NT2RP4001447	F-NT2RP4001447	0. 33
	NT2RP4001474	F-NT2RP4001474	0.89
20	NT2RP4001483	F-NT2RP4001483	0. 94
30	NT2RP4001498	F-NT2RP4001498	0. 59
	NT2RP4001502	F-NT2RP4001502	0. 61
	NT2RP4001507	F-NT2RP4001507	0. 91
	NT2RP4001524	F-NT2RP4001524	0.61
35	NT2RP4001529	F-NT2RP4001529	0.75
	NT2RP4001547	F-NT2RP4001547	0.77
	NT2RP4001551	F-NT2RP4001551	0.79
	NT2RP4001555	F-NT2RP4001555	0.79
40	NT2RP4001567	F-NT2RP4001567	0.39
	NT2RP4001568	F-NT2RP4001568	0.45
	NT2RP4001571	F-NT2RP4001571	0.76
	NT2RP4001574	F-NT2RP4001574	0.91
45	NT2RP4001575	F-NT2RP4001575	0.57
	NT2RP4001592	F-NT2RP4001592	0.71
	NT2RP4001610	F-NT2RP4001610	0. 74
	NT2RP4001614	F-NT2RP4001614	0. 73
50	NT2RP4001634	F-NT2RP4001634	0.73
	NT2RP4001638	F-NT2RP4001638	0. 40
	NT2RP4001644	F-NT2RP4001644	0. 94
	NT2RP4001656	F-NT2RP4001656	0. 92
55	NT2RP4001677	F-NT2RP4001677	0. 47
	NT2RP4001679	F-NT2RP4001679	0. 54

	NT2RP4001696	C HT0DD4004000	امن
		F-NT2RP4001696	0. 62
	NT2RP4001725	F-NT2RP4001725	0. 94
5	NT2RP4001730	F-NT2RP4001730	0. 94
5	NT2RP4001739	F-NT2RP4001739_	•
	NT2RP4001753	F-NT2RP4001753	0. 66
	NT2RP4001760	F-NT2RP4001760	0.39
	NT2RP4001790	F-NT2RP4001790	0.64
10	NT2RP4001803	F-NT2RP4001803	0.87
	NT2RP4001822	F-NT2RP4001822	0.84
	NT2RP4001823	F-NT2RP4001823	0.56
	NT2RP4001828	F-NT2RP4001828	0.45
15	NT2RP4001838	F-NT2RP4001838	0.87
	NT2RP4001841	F-NT2RP4001841	0.58
	NT2RP4001849	F-NT2RP4001849,	0.88
	NT2RP4001861	F-NT2RP4001861	0.80
20	NT2RP4001893	F-NT2RP4001893	0.94
	NT2RP4001896	F-NT2RP4001896	0. 93
	NT2RP4001901	F-NT2RP4001901	0. 89
	NT2RP4001927	F-NT2RP4001927	0.77
25	NT2RP4001938	F-NT2RP4001938	0.94
	NT2RP4001946	F-NT2RP4001946	0.74
	NT2RP4001950	F-NT2RP4001950	0.45
	NT2RP4001953	F-NT2RP4001953	0.49
30	NT2RP4001966	F-NT2RP4001966	0.86
	NT2RP4001975	F-NT2RP4001975	0.94
	NT2RP4002018	F-NT2RP4002018	0. 67
	NT2RP4002047	F-NT2RP4002047	0.67
35	NT2RP4002052	F-NT2RP4002052	0.42
	NT2RP4002058	F-NT2RP4002058	0.78
	NT2RP4002071	F-NT2RP4002071	0.94
	NT2RP4002078	F-NT2RP4002078	0.38
40	NT2RP4002081	F-NT2RP4002081	0.55
70	NT2RP4002083	F-NT2RP4002083	0.86
	NT2RP4002408	F-NT2RP4002408	0.64
	NT2RP4002791	F-NT2RP4002791	0.84
45	NT2RP4002888	F-NT2RP4002888	0.31
45	NT2RP5003459	F-NT2RP5003459	0.94
	NT2RP5003461	F-NT2RP5003461	0.94
	NT2RP5003477	F-NT2RP5003477	0.44
	NT2RP5003492	F-NT2RP5003492	0.57
50	NT2RP5003500	F-NT2RP5003500	0.94
	NT2RP5003506	F-NT2RP5003506	0.94
	NT2RP5003512	F-NT2RP5003512	0.42
	NT2RP5003522	F-NT2RP5003522	0.94
55	NT2RP5003524	F-NT2RP5003524	0.58
	NT2RP5003534	F-NT2RP5003534	0.32

	OVARC1000001	F-0VARC1000001	0.74
		F-0VARC1000004	0. 90
	0VARC1000006	F-0VARC1000004	0. 94
5	OVARC1000013	F-0VARC1000008	0. 48
	0VARC1000013	F-0VARC1000013	
			0. 69
	0VARC1000035	F-0VARC1000035	0. 59
10	0VARC1000060	F-0VARC1000060	0. 76
10	0VARC1000087	F-0VARC1000087	0. 31
	0VARC1000091	F-0VARC1000091	0. 88
	0VARC1000106	F-0VARC1000106	0. 88
	OYARC1000109	F-0VARC1000109	0. 94
15	0VARC1000113	F-0VARC1000113	0. 40
	OYARC1000114	F-0VARC1000114	0. 53
	OVARC1000139	F-0VARC1000139,	0. 88
	0VARC1000148	F-0VARC1000148	0. 94
20	OVARC1000151	F-0VARC1000151	0. 34
	0VARC1000168	F-0VARC1000168	0. 56
	0VARC1000209	F-0VARC1000209	0. 94
	0VARC1000212	F-0VARC1000212	0.87
25	OVARC1000241	F-0VARC1000241	0. 59
	OVARC1000288	F-0VARC1000288	0. 87
	OVARC1000304	F-0VARC1000304	0. 94
	0VARC1000309	F-0VARC1000309	0. 76
30	0VARC1000321	F-0VARC1000321	0. 90
	0VARC1000326	F-0VARC1000326	0. 46
	0VARC1000335	F-0VARC1000335	0. 55
	0VARC1000347	F-0VARC1000347	0. 43
35	0VARC1000384	F-0VARC1000384	0. 90
	0VARC1000408	F-0VARC1000408	0. 42
	0VARC1000411	F-0VARC1000411	0. 65
	0VARC1000420	F-0VARC1000420	0. 84
40	0VARC1000437	F-0VARC1000437	0. 94
40	0VARC1000440	F-0VARC1000440	0. 94
	0VARC1000442	F-0VARC1000442	0.94
	0VARC1000443	F-0VARC1000443	0.74
45	OVARC1000461	F-0VARC1000461	0.89
45	OVARC1000465	F-0VARC1000465	0. 75
	OVARC1000466	F-0VARC1000466	0.94
	0VARC1000473	F-0VARC1000473	0.31
	0VARC1000479	F-0VARC1000479	0.42
50	0VARC1000520	F-0VARC1000520	0. 94
	0VARC1000556	F-0VARC1000556	0.94
	OVARC1000557	F-0VARC1000557	0. 72
	0YARC1000564	F-0VARC1000564	0.94
55	0VARC1000576	F-0VARC1000576	0.40
	OVARC1000588	F-0YARC1000588	0. 45

	OVARC1000605	F-0VARC1000605	0. 58
	0VARC1000622	F-0VARC1000622	0.80
	OVARC1000640	F-0VARC1000640	0. 35
5	OVARC1000649	F-0VARC1000649	
	OVARC1000643	F-0VARC1000661	0.87
	OVARC1000681	F-0VARC1000681	0. 84
10	OVARC1000682	F-0VARC1000682	0.38
10	OVARC1000689	F-0VARC1000689	0.36
	OVARC1000703	F-0VARC1000703	0. 92
	OVARC1000722	F-0VARC1000722	0. 94
	OVARC1000730	F-0VARC1000730	0.90
15	OVARC1000746	F-0VARC1000746	0.42
	OVARC1000771		0.94
	OVARC1000781		i i
	OVARC1000787	F-0VARC1000787	0.41
20	0VARC1000800	F-0VARC1000800	0.85
	0VARC1000834	F-0VARC1000834	0.62
	0VARC1000846	F-0VARC1000846	0.49
	0VARC1000850	F-0VARC1000850	0.94
25	0VARC1000862	F-0VARC1000862	0. 54
•	0VARC1000876	F-0VARC1000876	0.74
	0VARC1000883	F-0VARC1000883	0. 55
	0VARC1000885	F-0VARC1000885	0.94
30	0VARC1000886	F-0VARC1000886	0.61
	0VARC1000890	F-0VARC1000890	0.45
	0VARC1000912	F-0VARC1000912	0.63
	0VARC1000915	F-0VARC1000915	0.31
35	0VARC1000924	F-0VARC1000924	0.82
	0VARC1000945	F-0VARC1000945	0.74
	0VARC1000959	F-0VARC1000959	0. 33
	0VARC1000964	F-0VARC1000964	0.60
40	OVARC1000984	F-0VARC1000984	0. 59
40	0VARC1000996	F-0VARC1000996	0.83
•	0VARC1000999	F-0VARC1000999	0.94
	0VARC1001004	F-0VARC1001004	0. 69
45	0VARC1001010	F-0VARC1001010	0.51
45	0VARC1001032	F-0VARC1001032	0. 32
	0VARC1001034	F-0VARC1001034	0.78
	0VARC1001038	F-0VARC1001038	0. 93
	0VARC1001040	F-0VARC1001040	0.68
50	0VARC1001044	F-0VARC1001044	0.66
	0VARC1001055	F-0VARC1001055	0. 76
	0VARC1001065	F-0VARC1001065	0.31
	0VARC1001068	F-0VARC1001068	0.63
55	0VARC1001074	F-0VARC1001074	0.82
	OVARC1001092	F-0VARC1001092	0.50

	OVARC1001107	F-0VARC1001107	0.65
	OVARC1001113	F-0VARC1001113	0.86
	0VARC1001154	F-0VARC1001154	0.43
5	0VARC1001161	F-0VARC1001161	
	OVARC1001162	F-0VARC1001162	0. 61
	OVARC1001167	F-0VARC1001167	0. 31
	OVARC1001170	F-0VARC1001170	0.46
10	OVARC1001171	F-0VARC1001171	0. 78
	OVARC1001173	F-0VARC1001173	0.71
	OVARC1001176	F-0VARC1001176	0. 83
	OVARC1001170		0. 53
	OVARC1001188	F-0VARC1001180	
15		F-0VARC1001188	0.42
	0VARC1001200	F-0VARC1001200	0.94
	OVARC1001232	F-0VARC1001232,	
	OVARC1001243	F-0VARC1001243	0. 59
20	0VARC1001244	F-0VARC1001244	0.79
	0VARC1001270	F-0VARC1001270	0. 48
	0VARC1001271	F-0VARC1001271	0. 67
	0VARC1001296		0, 60
25	OVARC1001306	F-0VARC1001306	0. 44
	0VARC1001329	F-0VARC1001329	0. 91
	OVARC1001341	F-0VARC1001341	0. 82
	OVARC1001344	F-0VARC1001344	0. 94
30	0VARC1001360	F-0VARC1001360	0. 94
30	OVARC1001369	F-0VARC1001369	0. 36
	OVARC1001372	F-0VARC1001372	0.63
	0VARC1001376	F-0VARC1001376	0.81
	OVARC1001381	F-0VARC1001381	0.78
35	OVARC1001391	F-0VARC1001391	0. 33
	OVARC1001399	F-0VARC1001399	0.67
	OVARC1001417	F-0VARC1001417	0. 94
	OVARC1001419	F-0VARC1001419	0. 34
40	0VARC1001425	F-0VARC1001425	0.73
	0VARC1001436	F-0VARC1001436	0. 59
	OVARC1001453	F-0VARC1001453	0. 45
	OVARC1001476	F-0VARC1001476	0.85
45	0VARC1001489	F-0VARC1001489	0.35
	0VARC1001496	F-0VARC1001496	0.90
	OVARC1001506	F-0VARC1001506	0. 59
	0VARC1001525	F-0VARC1001525	0. 87
50	0VARC1001555	F-0VARC1001555	0. 67
	0VARC1001577	F-0VARC1001577	0. 94
	OVARC1001610	F-0VARC1001610	0.38
	0VARC1001611	F-0VARC1001611	0.34
55	OVARC1001702	F-0VARC1001702	0.88
55	OVARC1001703	F-0VARC1001703	0. 76

	OVARC1001711	F-0VARC1001711	0. 82
	OVARC1001713	F-0VARC1001713	0. 82
	OVARC1001731	F-0VARC1001731	0. 94
5	OVARC1001762	F-0VARC1001762	0. 59
	OVARC1001766	F-0VARC1001766	0. 94
	OVARC1001767	F-0VARC1001767	0. 70
	OVARC1001768	F-0VARC1001768	0. 32
10	OVARC1001791	F-0VARC1001791	0. 74
	0VARC1001802	F-0VARC1001802	0. 76
	0VARC1001809	F-0VARC1001809	0. 63
	0VARC1001820	F-0VARC1001820	0. 94
15	OVARC1001828	F-0VARC1001828	0. 78
	OVARC1001861	F-0VARC1001861	0. 94
	OVARC1001873	F-0VARC1001873,	
	OVARC1001879	F-0VARC1001879	0. 94
20	OVARC1001880	F-0VARC1001880	0. 46
	OVARC1001916	F-0VARC1001916	0. 82
	OVARC1001942	F-0VARC1001942	0. 72
	OVARC1001943	F-0VARC1001943	0.49
25	OVARC1001949	F-0VARC1001949	0.46
	0VARC1001950	F-0VARC1001950	0. 94
	OVARC1001987	F-0VARC1001987	0.91
	0VARC1002050	F-0VARC1002050	0.85
30	OVARC1002082	F-0VARC1002082	0.94
	OVARC1002107	F-0VARC1002107	0.33
	OVARC1002112	F-0VARC1002112	0. 94
	OVARC1002127	F-0VARC1002127	0. 80
35	0VARC1002138	F-0VARC1002138	0. 46
	OVARC1002156	F-0VARC1002156	0. 43
	OVARC1002158	F-0VARC1002158	0. 94
	OVARC1002165	F-0VARC1002165	0. 88
40	OVARC1002182	F-0VARC1002182	0. 94
	PLACE1000004	F-PLACE1000004	0. 43
	PLACE1000005	F-PLACE1000005	0. 37
	PLACE1000007	F-PLACE1000007	0. 72
45	PLACE1000014	F-PLACE1000014	0. 38
	PLACE1000048	F-PLACE1000048	0. 36
	PLACE1000050	F-PLACE1000050	0. 94
	PLACE1000061	F-PLACE1000061	0. 56
50	PLACE1000066	F-PLACE1000066	0. 94
	PLACE1000081	F-PLACE1000081	0. 37
	PLACE1000133	F-PLACE1000133	0. 53
	PLACE1000142	F-PLACE1000142	0. 94
55	PLACE1000184	F-PLACE1000184	0. 73
	PLACE1000185	F-PLACE1000185	0.86
	PLACE1000213	F-PLACE1000213	0. 34

	PLACE1000236	F-PLACE1000236	0. 76
	PLACE1000246	F-PLACE1000246	0.91
	PLACE1000308	F-PLACE1000308	0.40
5	PLACE1000347	F-PLACE1000347	0.48
	PLACE1000374	F-PLACE1000374	0. 58
	PLACE1000380	F-PLACE1000380	0. 94
	PLACE1000383	F-PLACE1000383	0. 39
10	PLACE1000401	F-PLACE1000401	0. 94
	PLACE1000406	F-PLACE1000406	0. 94
·	PLACE1000420	F-PLACE1000420	0. 82
	PLACE1000435	F-PLACE1000435	0. 32
15	PLACE1000444	F-PLACE1000444	0. 41
	PLACE1000453	F-PLACE1000453	0. 76
	PLACE1000492	F-PLACE1000492,	0. 65
	PLACE1000547	F-PLACE1000547	0. 94
20	PLACE1000562	F-PLACE1000562	0. 94
20	PLACE1000562	F-PLACE1000564	0. 45
	PLACE1000588	F-PLACE1000588	0. 43
	PLACE1000596	F-PLACE1000596	0. 60
	PLACE1000596	F-PLACE1000596	0. 94
25	PLACE1000611	F-PLACE1000611	0. 48
	PLACE1000611	F-PLACE1000611	0. 40
	PLACE1000653	F-PLACE1000653	0. 92
	PLACE1000656	F-PLACE1000656	0. 94
30			0. 94
	PLACE1000706	F-PLACE1000706	
	PLACE1000712	F-PLACE1000712	0. 78 0. 94
	PLACE1000716	F-PLACE1000716	
35	PLACE1000748	F-PLACE1000748	0. 52
	PLACE1000755	F-PLACE1000755	0.87
	PLACE1000769	F-PLACE1000769	0. 94
	PLACE1000785	F-PLACE1000785	0. 33
40	PLACE1000786	F-PLACE1000786	0.88
	PLACE1000793	F-PLACE1000793	0. 79
	PLACE1000798	F-PLACE1000798	0.59
	PLACE1000849	F-PLACE1000849	0. 64
45	PLACE1000856	F-PLACE1000856	0.40
	PLACE1000863	F-PLACE1000863	0. 94
	PLACE1000909	F-PLACE1000909	0. 94
	PLACE1000948	F-PLACE1000948	0. 92
50	PLACE1000972	F-PLACE1000972	0. 73
	PLACE1000977	F-PLACE1000977	0. 72
	PLACE1000987	F-PLACE1000987	0. 93
	PLACE1001000	F-PLACE1001000	0. 37
55	PLACE1001010	F-PLACE1001010	0. 50
	PLACE1001024	F-PLACE1001024	0. 94
	PLACE1001036	F-PLACE1001036	0.51

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	PLACE1001054	F-PLACE1001054	0.88
	PLACE1001062	F-PLACE1001062	0.57
	PLACE1001092	F-PLACE1001092	0.94
5	PLACE1001104	F-PLACE1001104	0.38
	PLACE1001168	F-PLACE1001168	0.41
	PLACE1001171	F-PLACE1001171	0.36
	PLACE1001185	F-PLACE1001185	0. 93
10	PLACE1001238	F-PLACE1001238	0. 87
,,,	PLACE1001241	F~PLACE1001241	0. 67
	PLACE1001257	F-PLACE1001257	0.40
	PLACE1001280	F-PLACE1001280	0. 70
45	PLACE1001280	F-PLACE1001294	0. 70
15			
	PLACE1001304	F-PLACE1001304	0. 94
	PLACE1001311	F-PLACE1001311,	
	PLACE1001351	F-PLACE1001351	0.86
20	PLACE1001366	F-PLACE1001366	0. 39
	PLACE1001383	F-PLACE1001383	0.94
	PLACE1001387	F-PLACE1001387	0. 34
	PLACE1001399	F-PLACE1001399	0.38
25	PLACE1001412	F-PLACE1001412	0.94
	PLACE1001456	F-PLACE1001456	0.42
	PLACE1001484	F-PLACE1001484	0.88
	PLACE1001503	F-PLACE1001503	0.85
30	PLACE1001545	F-PLACE1001545	0.92
	PLACE1001551	F-PLACE1001551	0.39
	PLACE1001570	F-PLACE1001570	0. 40
	PLACE1001602	F-PLACE1001602	0.60
35	PLACE1001608	F-PLACE1001608	0.40
	PLACE1001610	F-PLACE1001610	0.43
	PLACE1001632	F-PLACE1001632	0.48
	PLACE1001634	F-PLACE1001634	0.60
40	PLACE1001692	F-PLACE1001692	0.74
40	PLACE1001729	F-PLACE1001729	0.69
	PLACE1001739	F-PLACE1001739	0.49
	PLACE1001740	F-PLACE1001740	0.50
	PLACE1001761	F-PLACE1001761	0.71
45	PLACE1001771	F-PLACE1001771	0.39
	PLACE1001781	F-PLACE1001781	0.57
	PLACE1001810	F-PLACE1001810	0.57
	PLACE1001817	F-PLACE1001817	0.91
50	PLACE1001844	F-PLACE1001844	0. 34
	PLACE1001869	F-PLACE1001869	0.34
	PLACE1001912	F-PLACE1001912	0.82
	PLACE1001920	F-PLACE1001920	0.85
55	PLACE1001928	F-PLACE1001928	0.60
	PLACE1001983	F-PLACE1001983	0.39
	. 3.02.001000	, , 2.021001000	5. 50

	PLACE1001989	F-PLACE1001989	0:74
	PLACE1002004	F-PLACE1002004	0.70
	PLACE1002046	F-PLACE1002046	0.81
5	PLACE1002072	F-PLACE1002072_	0. 94
	PLACE1002073	F-PLACE1002073	0. 70
	PLACE1002090	F-PLACE1002090	0.64
	PLACE1002115	F-PLACE1002115	0. 60
10	PLACE1002119	F-PLACE1002119	0.71
	PLACE1002140	F-PLACE1002140	0.94
	PLACE1002163	F-PLACE1002163	0. 56
	PLACE1002170	F-PLACE1002170	0.60
45	PLACE1002170	F-PLACE1002171	0.72
15	PLACE1002171	F-PLACE1002213	0. 72
	PLACE1002342	F-PLACE1002342,	
	PLACE1002395	F-PLACE1002395	0.87
20	PLACE1002433	F-PLACE1002433	0.73
	PLACE1002437	F-PLACE1002437	0.46
	PLACE1002438	F-PLACE1002438	0.41
	PLACE1002450	F-PLACE1002450	0.48
25	PLACE1002465	F-PLACE1002465	0. 34
	PLACE1002474	F-PLACE1002474	0.89
	PLACE1002499	F-PLACE1002499	0. 94
	PLACE1002529	F-PLACE1002529	0. 45
30	PLACE1002532	F-PLACE1002532	0. 76
,	PLACE1002571	F-PLACE1002571	0.40
	PLACE1002625	F-PLACE1002625	0. 36
	PLACE1002655	F-PLACE1002655	0.41
35	PLACE1002665	F-PLACE1002665	0.80
35	PLACE1002685	F-PLACE1002685	0. 31
	PLACE1002714	F-PLACE1002714	0.90
	PLACE1002722	F-PLACE1002722	0. 43
	PLACE1002775	F-PLACE1002775	0. 94
40	PLACE1002794	F-PLACE1002794	0. 90
	PLACE1002811	F-PLACE1002811	0.88
	PLACE1002815	F-PLACE1002815	0.40
	PLACE1002816	F-PLACE1002816	0.73
45	PLACE1002834	F-PLACE1002834	0. 41
	PLACE1002839	F-PLACE1002839	0.73
	PLACE1002851	F-PLACE1002851	0.90
	PLACE1002908	F-PLACE1002908	0.94
50	PLACE1002941	F-PLACE1002941	0. 33
	PLACE1002991	F-PLACE1002991	0.39
	PLACE1002993	F-PLACE1002993	0. 94
	PLACE1002996	F-PLACE1002996	0.40
55	PLACE1003025	F-PLACE1003025	0.77
	PLACE1003027	F-PLACE1003027	0.54

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	PLACE1003044	F-PLACE1003044	0.71
	PLACE1003045	F-PLACE1003045	0.89
	PLACE1003092	F-PLACE1003092	0. 59
5	PLACE1003100	F-PLACE1003100	0.38
	PLACE1003108	F-PLACE1003108	0.94
	PLACE1003136	F-PLACE1003136	0. 87
	PLACE1003145	F-PLACE1003145	0. 94
10	PLACE1003174	F-PLACE1003174	0.46
	PLACE1003176		0. 48
	PLACE1003190		0. 57
	PLACE1003200		0. 47
15	PLACE1003249		0. 33
,,	PLACE1003256		0. 57
	PLACE1003296		
	PLACE1003302	•	0. 94
20	PLACE1003334		0.78
20	PLACE1003342		0. 34
	PLACE1003353		0.71
	PLACE1003366		0. 66
	PLACE1003369		0. 91
25	PLACE1003383		0. 70
	PLACE1003394		0. 94
	PLACE1003454		0.94
	PLACE1003493		0.93
30	PLACE1003537	F-PLACE1003537	0.94
	PLACE1003553		0. 34
	PLACE1003592		0.33
	PLACE1003596	F-PLACE1003596	0.89
35	PLACE1003602	F-PLACE1003602	0.94
	PLACE1003605	F-PLACE1003605	0.94
	PLACE1003611	F-PLACE1003611	0.80
	PLACE1003625	F-PLACE1003625	0.39
40	PLACE1003669	F-PLACE1003669	0. 79
	PLACE1003704	F-PLACE1003704	0.43
	PLACE1003709	F-PLACE1003709	0. 90
	PLACE1003711	F-PLACE1003711	0.74
45	PLACE1003723	F-PLACE1003723	0.63
	PLACE1003738	F-PLACE1003738	0.91
	PLACE1003762	F-PLACE1003762	0.94
	PLACE1003771	F-PLACE1003771	0.56
50	PLACE1003783	F-PLACE1003783	0.39
	PLACE1003784	F-PLACE1003784	0.61
	PLACE1003795	F-PLACE1003795	0. 32
	PLACE1003870	F-PLACE1003870	0. 94
55	PLACE1003885	F-PLACE1003885	0. 94
	PLACE1003886	F-PLACE1003886	0. 94

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	PLACE1003888	F-PLACE1003888	0. 36
	PLACE1003892	F-PLACE1003892	0. 92
	PLACE1003903	F-PLACE1003903	0. 68
5	PLACE1003915	F-PLACE1003915	
	PLACE1003923	F-PLACE1003923	0. 52
	PLACE1003936	F-PLACE1003936	0. 67
	PLACE1003958	F-PLACE1003968	0. 59
10			
	PLACE1004103	F-PLACE1004103	0.94
	PLACE1004104	F-PLACE1004104	0. 44
	PLACE1004114	F-PLACE1004114	0. 91
	PLACE1004128	F-PLACE1004128	0. 72
15	PLACE1004149	F-PLACE1004149	0. 41
	PLACE1004156	F-PLACE1004156	0. 50
	PLACE1004161	F-PLACE1004161	0. 77
	PLACE1004183	F-PLACE1004183	0. 62
20	PLACE1004197	F-PLACE1004197	0.39
	PLACE1004203	F-PLACE1004203	0. 94
	PLACE1004258	F-PLACE1004258	0.46
	PLACE1004270	F-PLACE1004270	0. 65
25	PLACE1004277	F-PLACE1004277	0.86
	PLACE1004289	F-PLACE1004289	0. 79
	PLACE1004302	F-PLACE1004302	0. 57
	PLACE1004316	F-PLACE1004316	0. 64
30	PLACE1004358	F-PLACE1004358	0. 85
30	PLACE1004376	F-PLACE1004376	0. 64
	PLACE1004388	F-PLACE1004388	0.94
	PLACE1004405	F-PLACE1004405	0. 82
	PLACE1004428	F-PLACE1004428	0. 68
35	PLACE1004437	F-PLACE1004437	0.94
	PLACE1004451	F-PLACE1004451	0. 39
	PLACE1004460	F-PLACE1004460	0. 92
	PLACE1004473	F-PLACE1004473	0. 94
40	PLACE1004510	F-PLACE1004510	0. 94
	PLACE1004516	F-PLACE1004516	0.58
	PLACE1004548	F-PLACE1004548	0. 59
	PLACE1004550	F-PLACE1004550	0. 94
45	PLACE1004564	F-PLACE1004564	0. 53
	PLACE1004629	F-PLACE1004629	0. 34
		F-PLACE1004645	0. 59
	PLACE1004646	F-PLACE1004646	0.35
50	PLACE1004664	F-PLACE1004644	0. 33
	PLACE1004604	F-PLACE10046672	0. 83
	PLACE1004672	F-PLACE1004674	0. 90
			0. 90
	PLACE1004686	F-PLACE1004686	
55	PLACE1004691		0. 89
	PLACE1004722	F-PLACE1004722	0. 48

	PLACE1004736	F-PLACE1004736	0. 80
	PLACE1004740	F-PLACE1004740	0. 34
	PLACE1004743	F-PLACE1004743	0. 70
5	PLACE1004751	F-PLACE1004751	0.49
	PLACE1004777	F-PLACE1004777	0. 72
	PLACE1004804	F-PLACE1004804	0.46
	PLACE1004814	F-PLACE1004814	0.70
10	PLACE1004824	F-PLACE1004824	0.63
	PLACE1004868	F-PLACE1004868	0. 94
	PLACE1004885	F-PLACE1004885	0. 63
	PLACE1004902	F-PLACE1004902	0. 56
15	PLACE1004918	F-PLACE1004918	0. 85
	PLACE1004930	F-PLACE1004930	0.83
	PLACE1004937	F-PLACE1004937,	0.46
	PLACE1004969	F-PLACE1004969	0.62
20	PLACE1004982	F-PLACE1004982	0.61
	PLACE1005026	F-PLACE1005026	0.81
	PLACE1005027	F-PLACE1005027	0. 91
	PLACE1005046		0.31
25	PLACE1005055		0. 57
	PLACE1005066		0. 68
	PLACE1005101		0. 94
	PLACE1005102		0.94
30	PLACE1005181		0. 94
		F-PLACE1005187	0. 94
	PLACE1005206		0. 67
	PLACE1005232		0. 72
35	PLACE1005243 PLACE1005261	F-PLACE1005243 F-PLACE1005261	0. 81 0. 75
	PLACE1005261		0. 75
	PLACE1005277		0. 33
		F-PLACE1005287	0.77
40	PLACE1005305		0. 94
	PLACE1005308	· - · · · · · · · ·	0.46
	PLACE1005313		0. 94
•	PLACE1005327		0. 82
45	PLACE1005331	F-PLACE1005331	0. 94
	PLACE1005335	F-PLACE1005335	0. 94
	PLACE1005373	F-PLACE1005373	0.71
	PLACE1005374	F-PLACE1005374	0.82
50	PLACE1005480	F-PLACE1005480	0. 43
	PLACE1005481	F-PLACE1005481	0. 42
	PLACE1005494	F-PLACE1005494	0. 34
	PLACE1005530	F-PLACE1005530	0. 94
55	PLACE1005550	F-PLACE1005550	0. 86
	PLACE1005554	F-PLACE1005554	0. 34

	PLACE1005557	F-PLACE1005557	0. 80
	PLACE1005595	F-PLACE1005595	0. 69
	PLACE1005603	F-PLACE1005603	0. 92
5	PLACE1005623	F-PLACE1005623	
	PLACE1005630	F-PLACE1005630	0. 58
	PLACE1005646	F-PLACE1005646	0.31
	PLACE1005656	F-PLACE1005656	0.85
10	PLACE1005698	F-PLACE1005698	0. 91
	PLACE1005727	F-PLACE1005727	0. 94
	PLACE1005739	F-PLACE1005739	0.87
٧	PLACE1005755	F-PLACE1005755	0. 32
15	PLACE1005763	F-PLACE1005763	0. 74
	PLACE1005799	F-PLACE1005799	0. 47
	PLACE1005803	F-PLACE1005803,	
	PLACE1005804	F-PLACE1005804	0. 36
20	PLACE1005813	F-PLACE1005813	0.76
	PLACE1005828	F-PLACE1005828	0. 43
	PLACE1005834	F-PLACE1005834	0. 89
	PLACE1005851	F-PLACE1005851	0. 39
25	PLACE1005876	F-PLACE1005876	0.51
-	PLACE1005890	F-PLACE1005890	0.81
	PLACE1005921	F-PLACE1005921	0. 90
	PLACE1005923	F-PLACE1005923	0.70
30	PLACE1005925	F-PLACE1005925	0.55
	PLACE1005934	F-PLACE1005934	0.94
	PLACE1005936	F-PLACE1005936	0.41
	PLACE1005951	F-PLACE1005951	0.34
35	PLACE1005953	F-PLACE1005953	0.31
	PLACE1005955	F-PLACE1005955	0.89
	PLACE1005966	F-PLACE1005966	0.73
	PLACE1005990	F-PLACE1005990	0.37
40	PLACE1006011	F-PLACE1006011	0.94
	PLACE1006037	F-PLACE1006037	0.65
	PLACE1006040	F-PLACE1006040	0. 94
	PLACE1006119	F-PLACE1006119	0.52
45	PLACE1006139	F-PLACE1006139	0.94
40	PLACE1006157	F-PLACE1006157	0.34
	PLACE1006159	F-PLACE1006159	0.50
	PLACE1006167	F-PLACE1006167	
50	PLACE1006170	F-PLACE1006170	0.94
50	PLACE1006195	F-PLACE1006195	0.31
	PLACE1006196	F-PLACE1006196	0.84
	PLACE1006225	F-PLACE1006225	0.87
EE	PLACE1006236	F-PLACE1006236	0. 92
55	PLACE1006239	F-PLACE1006239	0. 53
	PLACE1006246	F-PLACE1006246	0. 44
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	PLACE1006248	F-PLACE1006248	0.49
	PLACE1006325	F-PLACE1006325	0.74
	PLACE1006335		0.82
5	PLACE1006357		
	PLACE1006385	** ***	0. 54
	PLACE1006412		0. 79
	PLACE1006414		0. 73
10			
10	PLACE1006438		0. 94
	PLACE1006445		0. 37
	PLACE1006469		0. 85
	PLACE1006482		0. 94
15	PLACE1006488		0.94
	PLACE1006492		0. 32
	PLACE1006531	•	0. 45
	PLACE1006552	F-PLACE1006552	0. 42
20	PLACE1006598	F-PLACE1006598	0. 33
	PLACE1006615	F-PLACE1006615	0. 89
	PLACE1006626	F-PLACE1006626	0.69
	PLACE1006673	F-PLACE1006673	0.61
25	PLACE1006678	F-PLACE1006678	0. 53
	PLACE1006704	F-PLACE1006704	0. 47
	PLACE1006731	F-PLACE1006731	0.72
	PLACE1006782	F-PLACE1006782	0. 68
30	PLACE1006819	F-PLACE1006819	0. 57
30	PLACE1006829	F-PLACE1006829	0. 47
	PLACE1006883	F-PLACE1006883	0. 88
	PLACE1006901	F-PLACE1006901	0. 43
	PLACE1006917	F-PLACE1006917	0. 56
35	PLACE1006932		0. 45
	PLACE1006935	F-PLACE1006935	0. 58
	PLACE1006956	F-PLACE1006956	0.77
	PLACE1006958	F-PLACE1006958	0, 75
40	PLACE1006961		0. 34
	PLACE1006962		0. 94
	PLACE1006966		0. 56
	PLACE1006989		0. 47
45	PLACE1007014		0. 94
	PLACE1007021	F-PLACE1007021	0. 74
	PLACE1007053	F-PLACE1007053	0. 73
	PLACE1007068	F-PLACE1007068	0. 43
50	PLACE1007008	F-PLACE1007105	0. 43
	PLACE1007103	F-PLACE1007105	0. 82
	PLACE1007112	F-PLACE1007178	0. 82
	PLACE1007178	F-PLACE1007178	0. 79
	PLACE1007226 PLACE1007238	F-PLACE1007228	0.79
55	PLACE1007238 PLACE1007239	F-PLACE1007238	
	FLAGE 100/239	F-FLAUE100/239	0.94

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	PLACE1007243	F-PLACE1007243	0.77
:	PLACE1007257	F-PLACE1007257	0. 54
5	PLACE1007274	F-PLACE1007274	
	PLACE1007282	F-PLACE1007282	0.94
	PLACE1007301	F-PLACE1007301	0.34
	PLACE1007317	F-PLACE1007317	0. 43
10	PLACE1007342	F-PLACE1007342	0. 72
	PLACE1007346	F-PLACE1007346	0. 92
	PLACE1007367	F-PLACE1007367	0.43
	PLACE1007375	F-PLACE1007375	0. 85
15	PLACE1007386	F-PLACE1007386	0. 58
	PLACE1007402	F-PLACE1007402	0.74
	PLACE1007409	F-PLACE1007409,	
	PLACE1007416	F-PLACE1007416	0. 80
20	PLACE1007450	F-PLACE1007450	0. 41
	PLACE1007452	F-PLACE1007452	0.68
	PLACE1007454	F-PLACE1007454	0.39
	PLACE1007460	F-PLACE1007460	0.58
25	PLACE1007484	F-PLACE1007484	0.94
20	PLACE1007488	F-PLACE1007488	0.50
	PLACE1007507	F-PLACE1007507	0.44
	PLACE1007511	F-PLACE1007511	0.86
30	PLACE1007524	F-PLACE1007524	0.66
30	PLACE1007537	F-PLACE1007537	0.93
	PLACE1007544	F-PLACE1007544	0.94
	PLACE1007547	F-PLACE1007547	0.65
35	PLACE1007583	F-PLACE1007583	0.87
	PLACE1007598	F-PLACE1007598	0.39
	PLACE1007618	F-PLACE1007618	0.62
	PLACE1007621	F-PLACE1007621	0. 63
40	PLACE1007632	F-PLACE1007632	0. 75
40	PLACE1007645	F-PLACE1007645	0. 94
	PLACE1007649	F-PLACE1007649	0. 56
	PLACE1007688	F-PLACE1007688	0. 94
45	PLACE1007690	F-PLACE1007690	0. 44
45	PLACE1007697	F-PLACE1007697	0. 93
	PLACE1007705	F-PLACE1007705	0.61
	PLACE1007706	F-PLACE1007706	0.47
	PLACE1007725	F-PLACE1007725	0.61
50	PLACE1007729	F-PLACE1007729	0. 53
	PLACE1007730	F-PLACE1007730	0. 31
	PLACE1007746	F-PLACE1007746	0.81
	PLACE1007791	F-PLACE1007791	0.46
55	PLACE1007810	F-PLACE1007810	0. 73
	PLACE1007843	F-PLACE1007843	0. 31

	PLACE1007846	F-PLACE1007846	0.38
	PLACE1007858	F-PLACE1007858	0. 79
	PLACE1007897		
5	•	F-PLACE1007897	0.94
3	PLACE1007908		0.94
	PLACE1007946	F-PLACE1007946	0.52
	PLACE1007954	F-PLACE1007954	0.43
	PLACE1007955	F-PLACE1007955	0.41
10	PLACE1007958	F-PLACE1007958	0.37
	PLACE1007969	F-PLACE1007969	0.87
	PLACE1007990	F-PLACE1007990	0. 94
	PLACE1008000	F-PLACE1008000	0. 94
15	PLACE1008002	F-PLACE1008002	0. 93
	PLACE1008044	F-PLACE1008044	0.72
	PLACE1008095	F-PLACE1008095,	0.94
	PLACE1008122	F-PLACE1008122	0.94
20	PLACE1008129	F-PLACE1008129	0.80
	PLACE1008132	F-PLACE1008132	0. 37
	PLACE1008177	F-PLACE1008177	0.74
	PLACE1008198	F-PLACE1008198	0.94
25	PLACE1008209	F-PLACE1008209	0.54
	PLACE1008273	F-PLACE1008273	0.94
	PLACE1008275	F-PLACE1008275	0.82
	PLACE1008280	F-PLACE1008280	0.34
30	PLACE1008309	F-PLACE1008309	0. 90
30	PLACE1008329	F-PLACE1008329	0.38
	PLACE1008356	F-PLACE1008356	0.93
	PLACE1008368	F-PLACE1008368	0.43
	PLACE1008398	F-PLACE1008398	0.78
35	PLACE1008401	F-PLACE1008401	0.81
	PLACE1008402	F-PLACE1008402	0.88
	PLACE1008429	F-PLACE1008429	0.39
	PLACE1008457	F-PLACE1008457	0. 77
40	PLACE1008465	F-PLACE1008465	0.94
	PLACE1008488	F-PLACE1008488	0.42
	PLACE1008524	F-PLACE1008524	0.41
	PLACE1008531	F-PLACE1008531	0.94
45	PLACE1008532	F-PLACE1008532	0.61
	PLACE1008533	F-PLACE1008533	0.45
	PLACE1008568	F-PLACE1008568	0.65
	PLACE1008603	F-PLACE1008603	0.42
50	PLACE1008626	F-PLACE1008626	0. 94
	PLACE1008627	F-PLACE1008627	0. 72
	PLACE1008629	F-PLACE1008629	0. 33
	PLACE1008650	F-PLACE1008650	0. 79
55	PLACE1008693	F-PLACE1008693	0. 32
	PLACE1008696	F-PLACE1008696	0.39

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	PLACE1008790	F-PLACE1008790	0. 81
	PLACE1008808	F-PLACE1008808	0.51
	PLACE1008813	F-PLACE1008813	0. 36
5	PLACE1008854	F-PLACE1008854	0. 81
	PLACE1008867	F-PLACE1008867	0. 34
	PLACE1008887		0.49
	PLACE1008902		0. 91
10	PLACE1008925		0. 94
	PLACE1008934		0.70
	PLACE1009027	F-PLACE1009027	0. 32
	PLACE1009045	F-PLACE1009045	0. 42
15	PLACE1009060	F-PLACE1009060	0.69
	PLACE1009090	F-PLACE1009090	0, 46
	PLACE1009091	F-PLACE1009091,	
	PLACE1009094	F-PLACE1009094	0. 44
20	PLACE1009099		0. 56
20	PLACE1009110		0.84
	PLACE1009111		0.80
	PLACE1009113		0.88
25	PLACE1009130		0.60
25	PLACE1009158		0.41
	PLACE1009166		0. 94
	PLACE1009174		0.38
•	PLACE1009186		0.59
30	PLACE1009190	F-PLACE1009190	0.66
	PLACE1009230	F-PLACE1009230	0.87
	PLACE1009298	F-PLACE1009298	0.41
	PLACE1009319	F-PLACE1009319	0.53
35	PLACE1009328	F-PLACE1009328	0.66
	PLACE 1009335	F-PLACE1009335	0. 47
	PLACE1009338	F-PLACE1009338	0.70
	PLACE1009368	F-PLACE1009368	0.94
40	PLACE1009375	F-PLACE1009375	0.76
	PLACE1009388	F-PLACE1009388	0.46
	PLACE1009404	F-PLACE1009404	0. 72
	PLACE1009434	F-PLACE1009434	0. 50
45	PLACE1009443	F-PLACE1009443	0. 90
	PLACE1009444	F-PLACE1009444	0.33
	PLACE1009459	F-PLACE1009459	0. 94
	PLACE1009468	F-PLACE1009468	0.88
50	PLACE1009476	F-PLACE1009476	0.84
	PLACE1009524	F-PLACE1009524	0. 88
	PLACE1009542	F-PLACE1009542	0. 55
	PLACE1009571	F-PLACE1009571	0.61
55	PLACE1009581	F-PLACE1009581	0.94
	PLACE1009596	F-PLACE1009596	0. 44

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	PLACE1009607	F-PLACE1009607	0. 47
	PLACE1009622	F-PLACE1009622	0.65
	PLACE1009659	F-PLACE1009659	0.36
5	PLACE1009665	F-PLACE1009665	0.43
	PLACE1009670	F-PLACE1009670	0.33
	PLACE1009708	F-PLACE1009708	0.80
	PLACE1009721	F-PLACE1009721	0. 81
10	PLACE1009731	F-PLACE1009731	0. 47
	PLACE1009763	F-PLACE1009763	0. 94
	PLACE1009794	F-PLACE1009794	0. 33
	PLACE1009798	F-PLACE1009798	0. 34
15	PLACE1009845	F-PLACE1009845	0. 66
,,,	PLACE1009861	F-PLACE1009861	0. 72
	PLACE1009886	F-PLACE1009886,	
	PLACE1009908	F-PLACE1009908	0. 92
20	PLACE1009921	F-PLACE1009921	0.70
20	PLACE1009971	F-PLACE1009971	0.40
	PLACE1009992	F-PLACE1009992	0.89
	PLACE1009995	F-PLACE1009995	0.88
0.5	PLACE1009997	F-PLACE1009997	0.41
25	PLACE1010023	F-PLACE1010023	0.90
	PLACE1010031	F-PLACE1010031	0.71
	PLACE1010053	F-PLACE1010053	0.51
	PLACE1010074		0. 75
30	PLACE1010076	F-PLACE1010076	0. 53
	PLACE1010096	F-PLACE1010096	0. 44
	PLACE1010102	F-PLACE1010102	0. 77
	PLACE1010105	F-PLACE1010105	0. 58
35	PLACE1010106	F-PLACE1010106	0. 39
	PLACE1010134	F-PLACE1010134	0. 85
	PLACE1010148	F-PLACE1010148	0. 32
	PLACE1010152	F-PLACE1010152	0. 94
40	PLACE1010194	F-PLACE1010194	0. 94
	PLACE1010202	F-PLACE1010202	0. 33
	PLACE1010261	F-PLACE1010261	0.36
	PLACE1010274	F-PLACE1010274	0. 94
45	PLACE1010293	F-PLACE1010293	0. 55
	PLACE1010310	F-PLACE1010310	0. 41
	PLACE1010321	F-PLACE1010321	0.40
	PLACE1010324	F-PLACE1010324	0. 40
50	PLACE1010329	F-PLACE1010329	0. 36
	PLACE1010362	F-PLACE1010362	0. 34
	PLACE1010364	F-PLACE1010364	0. 37
	PLACE1010383	F-PLACE1010383	0.94
55	PLACE1010401	F-PLACE1010401	0. 53
	PLACE1010481	F-PLACE1010481	0. 74

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		OF TOTAL	
	PLACE1010491	F-PLACE1010491	0.31
	PLACE1010492	F-PLACE1010492	0.39
	PLACE1010522	F-PLACE1010522	0.92
5	PLACE1010529	F-PLACE1010529	
	PLACE1010547	F-PLACE1010547	0. 82
	PLACE1010580	F-PLACE1010580	0.72
	PLACE1010599	F-PLACE1010599	0. 83
40	PLACE1010599	F-PLACE1010616	0. 54
10	PLACE1010618	F-PLACE1010622	
			0. 92
	PLACE1010629	F-PLACE1010629	0.94
	PLACE1010630	F-PLACE1010630	0. 37
15	PLACE1010661	F-PLACE1010661	0. 56
	PLACE1010714	F-PLACE1010714	0. 68
	PLACE1010720	F-PLACE1010720,	
	PLACE1010743	F-PLACE1010743	0.75
20	PLACE1010761	F-PLACE1010761	0.69
	PLACE1010771	F-PLACE1010771	0.94
	PLACE1010786	F-PLACE1010786	0.80
	PLACE1010800	F-PLACE1010800	0.41
25	PLACE1010811	F-PLACE1010811	0.94
20	PLACE1010833	F-PLACE1010833	0.94
	PLACE1010856	F-PLACE1010856	0.37
	PLACE1010857	F-PLACE1010857	0.54
	PLACE1010870	F-PLACE1010870	0.50
30	PLACE1010877	F-PLACE1010877	0.77
	PLACE1010900	F-PLACE1010900	0. 70
	PLACE1010917	F-PLACE1010917	0. 94
	PLACE1010925	F-PLACE1010925	0. 79
35	PLACE1010926	F-PLACE1010926	0. 94
	PLACE1010942	F-PLACE1010942	0. 36
	PLACE1010944	F-PLACE1010944	0. 61
	PLACE1010954	F-PLACE1010954	0. 83
40	PLACE1010960	F-PLACE1010960	0. 38
	PLACE1010965	F-PLACE1010965	0. 68
•	PLACE1011026	F-PLACE1011026	0. 38
	PLACE1011041	F-PLACE1011041	0. 94
45	PLACE1011046	F-PLACE1011046	0. 85
40	PLACE1011054	F-PLACE1011054	0. 65
	PLACE1011056		
	PLACE1011090	F-PLACE1011090	0. 94
	PLACE1011114	F-PLACE1011114	0.94
50			
	PLACE1011160	F-PLACE1011160	0.62
	PLACE1011214	F-PLACE1011214	0.40
	PLACE1011219	F-PLACE1011219	0.48
55	PLACE1011221	F-PLACE1011221	0.94
	PLACE1011229	F-PLACE1011229	0. 46

PLACE1011263 F-PLACE1011263 PLACE1011273 F-PLACE1011273 PLACE1011291 F-PLACE1011291 PLACE1011310 F-PLACE1011310 PLACE1011325 F-PLACE1011325 PLACE1011332 F-PLACE1011332	0. 94 0. 32
PLACE1011291 F-PLACE1011291 PLACE1011310 F-PLACE1011310 PLACE1011325 F-PLACE1011325	0.32
5 PLACE1011310 F-PLACE1011310 PLACE1011325 F-PLACE1011325	
PLACE1011325 F-PLACE1011325	0.55
PLACE1011325 F-PLACE1011325	0.74
	0.39
/=/////	0.74
PLACE1011371 F-PLACE1011371	0. 32
10 PLACE1011399 F-PLACE1011399	0.89
PLACE1011433 F-PLACE1011433	0. 33
PLACE1011472 F-PLACE1011472	0. 59
PLACE1011477 F-PLACE1011477	0.80
	0.83
	0.82
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PLACE1011635 F-PLACE1011635,	- 1
PLACE1011646 F-PLACE1011646	0.37
20 PLACE1011664 F-PLACE1011664	0.65
PLACE1011675 F-PLACE1011675	0.89
PLACE1011682 F-PLACE1011682	0.81
PLACE1011725 F-PLACE1011725	0.86
25 PLACE1011783 F-PLACE1011783	0.78
PLACE1011858 F-PLACE1011858	0.76
PLACE1011875 F-PLACE1011875	0.49
PLACE1011891 F-PLACE1011891	0. 34
30 PLACE1011896 F-PLACE1011896	0. 34
PLACE1011922 F-PLACE1011922	0.46
PLACE1011923 F-PLACE1011923	0. 74
PLACE1011982 F-PLACE1011982	0.71
PLACE1012031 F-PLACE1012031	0.80
35 PLACE2000006 F-PLACE2000006	0. 61
PLACE2000007 F-PLACE2000007	0. 37
PLACE2000014 F-PLACE2000014	0. 67
PLACE2000015 F-PLACE2000015	0.53
40 PLACE2000021 F-PLACE2000021	0.36
PLACE2000030 F-PLACE2000030	0.83
FLACEZUUUGGU 1 - FLACEZUUUGGU	0. 44
PLACE2000030 F-PLACE2000030 PLACE2000033 F-PLACE2000033	0.88
PLACE2000033 F-PLACE2000033	0. 67
PLACE2000033 F-PLACE2000033 PLACE2000034 F-PLACE2000034	
PLACE2000033 F-PLACE2000034 PLACE2000034 F-PLACE2000039 F-PLACE2000039	0. 67
PLACE2000033 F-PLACE2000034 PLACE2000034 F-PLACE2000034 45 PLACE2000039 F-PLACE2000039 PLACE2000047 F-PLACE2000047	0. 67 0. 84
PLACE2000033 F-PLACE2000034 PLACE2000034 F-PLACE2000034 PLACE2000039 F-PLACE2000039 PLACE2000047 F-PLACE2000050	0. 67 0. 84 0. 41
PLACE2000033 F-PLACE2000034 PLACE2000034 F-PLACE2000034 PLACE2000039 F-PLACE2000039 PLACE2000047 F-PLACE2000047 PLACE2000050 F-PLACE2000050 PLACE2000062 F-PLACE2000062	0. 67 0. 84 0. 41 0. 94
PLACE2000033 F-PLACE2000034 PLACE2000034 F-PLACE2000034 PLACE2000039 F-PLACE2000039 PLACE2000047 F-PLACE2000047 PLACE2000050 F-PLACE2000050 PLACE2000062 F-PLACE2000062 F-PLACE2000072 F-PLACE2000072	0. 67 0. 84 0. 41 0. 94 0. 86
PLACE2000033 F-PLACE2000034 PLACE2000034 F-PLACE2000034 PLACE2000039 F-PLACE2000039 PLACE2000047 F-PLACE2000047 PLACE2000050 F-PLACE2000050 PLACE2000062 F-PLACE2000062 PLACE2000072 F-PLACE2000072 PLACE2000097 F-PLACE2000097 PLACE2000100 F-PLACE2000100	0. 67 0. 84 0. 41 0. 94 0. 86 0. 94 0. 75
PLACE2000033 F-PLACE2000034 PLACE2000034 F-PLACE2000034 PLACE2000039 F-PLACE2000039 PLACE2000047 F-PLACE2000047 PLACE2000062 F-PLACE2000062 PLACE2000072 F-PLACE2000072 PLACE2000097 F-PLACE2000097 PLACE2000110 F-PLACE2000111 PLACE2000111 F-PLACE2000111	0. 67 0. 84 0. 41 0. 94 0. 86 0. 94
PLACE2000033 F-PLACE2000034 PLACE2000034 F-PLACE2000034 PLACE2000039 F-PLACE2000039 PLACE2000047 F-PLACE2000047 PLACE2000050 F-PLACE2000050 PLACE2000062 F-PLACE2000062 PLACE2000072 F-PLACE2000072 PLACE2000097 F-PLACE2000097 PLACE2000100 F-PLACE2000100 PLACE2000111 F-PLACE2000111	0. 67 0. 84 0. 41 0. 94 0. 86 0. 94 0. 75 0. 62

	PLACE2000170	E_DI A0E0000170	0.00
			0.60
	PLACE2000172 PLACE2000223		0.55
E		F-PLACE2000223	0.54
5	PLACE2000235		•
	PLACE2000246		
	PLACE2000274	F-PLACE2000274	0.82
	PLACE2000302	F-PLACE2000302	0.92
10	PLACE2000341	F-PLACE2000341	0. 58
	PLACE2000359	F-PLACE2000359	0. 33
	PLACE2000371		0. 71
	PLACE2000373	F-PLACE2000373	0. 65
15	PLACE2000399	F-PLACE2000399	0. 41
	PLACE2000404	F-PLACE2000404	0. 94
	PLACE2000411	F-PLACE2000411,	
	PLACE2000427	F-PLACE2000427	0.60
20	PLACE2000433	F-PLACE2000433	0.61
	PLACE2000438	F-PLACE2000438	0. 56
	PLACE2000458	F-PLACE2000458	0. 82
	PLACE3000009		0. 56
25	PLACE3000020	F-PLACE3000020	0. 94
	PLACE3000029	F-PLACE3000029	0. 94
	PLACE3000059	F-PLACE3000059	0.51
	PLACE3000070	F-PLACE3000070	0. 67
30	PLACE3000103	F-PLACE3000103	0. 34
	PLACE3000121		0. 32
	PLACE3000142		0. 67
	PLACE3000145		0.84
35	PLACE3000147		0. 57
	PLACE3000148		0.94
	PLACE3000155	F-PLACE3000155	0. 42
	PLACE3000156	F-PLACE3000156	0.40
40	PLACE3000157	F-PLACE3000157	0.75
	PLACE3000160	F-PLACE3000160	0.44
	PLACE3000194	F-PLACE3000194	0.37
	PLACE3000197 PLACE3000221	F-PLACE3000197 F-PLACE3000221	0. 37 0. 49
45	PLACE3000221		0. 49
70	PLACE3000242	F-PLACE3000226	0. 39
	PLACE3000242	F-PLACE3000242	0. 56
	PLACE3000244 PLACE3000254	F-PLACE3000254	0.60
50	PLACE3000254	F-PLACE3000294	0. 42
30	PLACE3000304	F-PLACE3000310	0. 42
	PLACE3000310	F-PLACE3000310	0. 77
	PLACE3000320	F-PLACE3000320	0. 46
	PLACE3000321	F-PLACE3000321	0. 40
55	PLACE3000331	F-PLACE3000331	0. 45
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	PLACE3000350	F-PLACE3000350	0. 52
	PLACE3000352	F-PLACE3000352	0. 65
	PLACE3000353	F-PLACE3000353	0.71
5	PLACE3000363	F-PLACE3000363	0. 45
	PLACE3000405	F-PLACE3000405	0. 52
	PLACE3000413	F-PLACE3000413	0. 85
	PLACE3000416	F-PLACE3000416	0. 94
10	PLACE3000425		0.40
·	PLACE3000455		0. 45
	PLACE3000477	F-PLACE3000477	0. 79
	PLACE4000009		0.74
15	PLACE4000014	F-PLACE4000014	0. 46
7.5	PLACE4000014	F-PLACE4000014	0. 40
	PLACE4000049	,	
	PLACE4000052		0.80
20	PLACE4000063		0.69
•	PLACE4000089		0.44
	PLACE4000106	F-PLACE4000106	0. 34
	PLACE4000128		0. 89
25	PLACE4000129		0. 61
	PLACE4000131	F-PLACE4000131	0. 55
	PLACE4000147		0. 94
	PLACE4000156		0. 37
30	PLACE4000192	F-PLACE4000192	0.88
	PLACE4000211	F-PLACE4000211	0. 64
	PLACE4000222	F-PLACE4000222	0. 40
	PLACE4000230	F-PLACE4000230	0. 49
35	PLACE4000259	F-PLACE4000259	0. 84
30	PLACE4000261	F-PLACE4000261	0. 42
	PLACE4000269	F-PLACE4000269	0.60
	PLACE4000270	F-PLACE4000270	0.61
	PLACE4000300	F-PLACE4000300	0. 85
40	PLACE4000323	F-PLACE4000323	0.76
	PLACE4000326	F-PLACE4000326	0.86
	PLACE4000369	F-PLACE4000369	0. 87
	PLACE4000387	F-PLACE4000387	0. 41
45	PLACE4000392	F-PLACE4000392	0.39
	PLACE4000431	F-PLACE4000431	0.88
	PLACE4000445	F-PLACE4000445	0.40
	PLACE4000450	F-PLACE4000450	0. 89
50	PLACE4000465	F-PLACE4000465	0. 38
	PLACE4000487	F-PLACE4000487	0. 44
	PLACE4000489	F-PLACE4000489	0. 51
	PLACE4000521	F-PLACE4000521	0. 38
55	PLACE4000522	F-PLACE4000522	0. 33
	PLACE4000558	F-PLACE4000558	0. 44

	PLACE4000581	EDI ACE 4000E01	A EA
	PLACE4000590	F-PLACE4000581	0. 59
	PLACE4000593	F-PLACE4000590	0.38
5	-	F-PLACE4000593	0.33
•	PLACE4000612	F-PLACE4000612	
	PLACE4000638	F-PLACE4000638	0.38
	PLACE4000650	F-PLACE4000650	0.59
10	PLACE4000654	F-PLACE4000654	0. 57
	PLACE4000670	F-PLACE4000670	0. 71
	SKNMC1000011	F-SKNMC1000011	0.94
	SKNMC1000013	F-SKNMC1000013	0.66
	SKNMC1000046	F-SKNMC1000046	0. 72
15	SKNMC1000050	F-SKNMC1000050	0. 47
	SKNMC1000091	F-SKNMC1000091	0.68
	THYR01000017	F-THYR01000017,	
	THYR01000034	F-THYR01000034	0. 48
20	THYR01000040	F-THYR01000040	0. 32
	THYR01000070	F-THYR01000070	0. 75
	THYR01000072	F-THYR01000072	0. 94
	THYR01000107	F-THYR01000107	0. 94
25	THYR01000121	F-THYR01000121	0.46
	THYR01000124	F-THYR01000124	0. 37
	THYR01000163	F-THYR01000163	0.45
	THYR01000173	F-THYR01000173	0. 94
30	THYR01000197	F-THYR01000197	0. 78
	THYR01000199	F-THYR01000199	0. 55
	THYR01000206	F-THYR01000206	0. 59
	THYR01000242	F-THYR01000242	0. 69
35	THYR01000253	F-THYR01000253	0. 48
	THYR01000270	F-THYR01000270	0.85
	THYR01000288	F-THYR01000288	0.31
	THYR01000320	F-THYR01000320	0.94
40	THYR01000327	F-THYR01000327	0. 44
	THYR01000343	F-THYR01000343	0. 82
	THYR01000358	F-THYR01000358	0.89
	THYR01000368	F-THYR01000368	0. 94
45	THYR01000381	F-THYR01000381	0. 64
70	THYR01000387	F-THYR01000387	0. 57
	THYR01000394	F-THYR01000394	0. 94
	THYR01000395	F-THYR01000395	0. 91
50	THYR01000401	F-THYR01000401	0. 73
50	THYR01000488	F-THYR01000488	0. 94
	THYR01000501	F-THYR01000501	0. 74
	THYR01000502	F-THYR01000502	0. 64
55	THYR01000505	F-THYR01000505	0. 47
55	THYR01000569	F-THYR01000569	0. 76
	THYR01000570	F-THYR01000570	0. 79

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	THYR01000585	F-THYR01000585	0.60
	THYR01000605	F-THYR01000605	0.79
	THYR01000637	F-THYR01000637	0.80
5	THYR01000662	F-THYR01000662	0.77
	THYR01000666	F-THYR01000666	0. 47
	THYR01000676	F-THYR01000676	0. 94
	THYR01000684	F-THYR01000684	0. 72
10			
10	THYR01000712	F-THYR01000712	0.94
	THYR01000715	F-THYR01000715	0.94
	THYR01000734	F-THYR01000734	0.44
	THYR01000756	F-THYR01000756	0.94
15	THYR01000777	F-THYR01000777	0.73
	THYR01000783	F-THYR01000783	0.74
	THYR01000787	F-THYR01000787,	0.67
	THYR01000829	F-THYR01000829	0.43
20	THYR01000855	F-THYR01000855	0. 91
	THYR01000916	F-THYR01000916	0. 64
	THYR01000926	F-THYR01000926	0. 39
	THYR01000934	F-THYR01000934	0.94
25	THYR01000951	F-THYR01000951	0.94
20	THYR01000952	F-THYR01000952	0.67
	THYR01000983	F-THYR01000983	0.87
	THYR01000984	F-THYR01000984	0.94
00	THYR01000988	F-THYR01000988	0. 37
30	THYR01001033	F-THYR01001033	0.74
	THYR01001100	F-THYR01001100	0.74
	THYR01001120	F-THYR01001120	0. 87
	THYR01001134	F-THYR01001134	0. 93
35	THYR01001142	F-THYR01001142	0.61
	THYR01001173	F-THYR01001173	0.50
	THYR01001189	F-THYR01001189	0.40
	THYR01001204	F-THYR01001204	0. 94
40	THYR01001213	F-THYR01001213	0.94
	THYR01001271	F-THYR01001271	0.93
	THYRO1001287	F-THYR01001287	0. 32
	THYR01001313	F-THYR01001313	0. 53
45	THYRO1001320	F-THYR01001320	0, 68
	THYR01001347	F-THYR01001347	0. 68
	THYR01001363	F-THYR01001363	0. 39
	THYR01001374	F-THYR01001374	0. 94
50	THYR01001374	F-THYR01001403	0. 49
30	THYR01001405	F-THYR01001405	0. 90
	THYR01001406	F-THYR01001406	0. 90
	THYR01001458	F-THYR01001458	0. 94
		F-THYR01001458	0. 94
55	THYR01001487		l l
	THYR01001537	F-THYR01001537	0. 94
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	THYR01001584	F-THYR01001584	0.77
	THYR01001602	F-THYR01001602	0.64
	THYR01001605	F-THYR01001605	0. 52
5	THYR01001637	F-THYR01001637_	0. 68
•	THYR01001656	F-THYR01001656	0.82
	THYR01001661	F-THYR01001661	0.94
	THYR01001671	F-THYR01001671	0.94
		F-THYR01001703	0.87
10	THYR01001703		
	THYR01001721	F-THYR01001721	0.38
	THYR01001738	F-THYR01001738	0.51
	THYR01001793	F-THYR01001793	0.84
15	THYR01001809	F-THYR01001809	0.73
	THYR01001828	F-THYR01001828	0.88
	THYR01001854	F-THYR01001854,	0, 47
	VESEN1000122	F-VESEN1000122	0.36
20	Y79AA1000013	F-Y79AA1000013	0.94
20	Y79AA1000037	F-Y79AA1000037	0.94
	Y79AA1000059	F-Y79AA1000059	0.94
	Y79AA1000131	F-Y79AA1000131	0.56
	Y79AA1000181	F-Y79AA1000181	0.59
25	Y79AA1000202	F-Y79AA1000202	0, 45
	Y79AA1000214	F-Y79AA1000214	0.93
	Y79AA1000230	F-Y79AA1000230	0. 94
	Y79AA1000231	F-Y79AA1000231	0.36
30	Y79AA1000258	F-Y79AA1000258	0.36
	Y79AA1000268	F-Y79AA1000268	0.92
	Y79AA1000313	F-Y79AA1000313	0.91
	Y79AA1000328	F-Y79AA1000328	0.94
35	Y79AA1000342	F-Y79AA1000342	0.83
	Y79AA1000349	F-Y79AA1000349	0.50
	Y79AA1000355	F-Y79AA1000355	0, 94
	Y79AA1000368	F-Y79AA1000368	0. 91
40	Y79AA1000420	F-Y79AA1000420	0. 94
40	Y79AA1000469	F-Y79AA1000469	0.51
	Y79AA1000480	F-Y79AA1000480	0. 94
	Y79AA1000539	F-Y79AA1000539	0. 74
	Y79AA1000540	F-Y79AA1000540	0. 74
45	Y79AA1000560	F-Y79AA1000560	0. 93
	Y79AA1000574	F-Y79AA1000574	0.87
	Y79AA1000589	F-Y79AA1000589	0.94
50	Y79AA1000627	F-Y79AA1000627	0. 55
	Y79AA1000705	F-Y79AA1000705	0.76
	Y79AA1000734	F-Y79AA1000734	0.60
	Y79AA1000748		0. 45
55	Y79AA1000752		0.94
55	Y79AA1000774	F-Y79AA1000774	0. 38

	Y79AA1000782	F-Y79AA1000782	0. 47
	Y79AA1000784	F-Y79AA1000784	0. 93
•	Y79AA1000794	F-Y79AA1000794	0. 67
5	Y79AA1000800	F-Y79AA10008Q0	
	Y79AA1000824	F-Y79AA1000824	0.90
	Y79AA1000827	F-Y79AA1000827	0. 94
	Y79AA1000827	F-Y79AA1000827	0. 34
10	Y79AA1000850	F-Y79AA1000850	0. 92
	Y79AA1000962	F-Y79AA1000850	0. 69
	Y79AA1000962	F-Y79AA1000962	
	Y79AA1000968	F-Y79AA1000968	0.64
15	Y79AA1000908	F-Y79AA1000968	0, 64
75			0.66
	Y79AA1000985	F-Y79AA1000985	0. 66
	Y79AA1001023	F-Y79AA1001023	
•	Y79AA1001041	F-Y79AA1001041	0. 90
20	Y79AA1001048	F-Y79AA1001048	0. 81
	Y79AA1001068	F-Y79AA1001068	0, 64
	Y79AA1001077	F-Y79AA1001077	0. 78
	Y79AA1001078 Y79AA1001105	F-Y79AA1001078	0. 92
25	Y79AA1001105	F-Y79AA1001105	0.70
		F-Y79AA1001145	0.46
	Y79AA1001177	F-Y79AA1001177	0.74
	Y79AA1001185	F-Y79AA1001185	0. 74
30	Y79AA1001211	F-Y79AA1001211	0.94
•	Y79AA1001216 Y79AA1001228	F-Y79AA1001216 F-Y79AA1001228	0. 61 0. 56
	Y79AA1001233	F-Y79AA1001233	0. 83
	Y79AA1001235	F-Y79AA1001236	0. 83
35	Y79AA1001281	F-Y79AA1001281	0. 52
	Y79AA1001299	F-Y79AA1001299	0. 94
	Y79AA1001312	F-Y79AA1001312	0. 53
	Y79AA1001312	F-Y79AA1001323	0. 76
40	Y79AA1001323	F-Y79AA1001391	0. 70
	Y79AA1001394	F-Y79AA1001394	0. 94
	Y79AA1001402	F-Y79AA1001402	0. 90
	Y79AA1001493	F-Y79AA1001493	0.94
45	Y79AA1001511	F-Y79AA1001511	0. 47
40	Y79AA1001533	F-Y79AA1001533	0. 92
	Y79AA1001541	F-Y79AA1001541	0. 32
	Y79AA1001548	F-Y79AA1001548	0.50
	Y79AA1001555	F~Y79AA1001555	0. 92
50	Y79AA1001581	F-Y79AA1001581	0. 94
	Y79AA1001585	F-Y79AA1001585	0.80
	Y79AA1001603	F-Y79AA1001603	0. 94
	Y79AA1001613	F-Y79AA1001613	0. 94
55	Y79AA1001647	F-Y79AA1001647	0. 49

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	Y79AA1001665	F-Y79AA1001665	0.78
	Y79AA1001679	F-Y79AA1001679	0.94
	Y79AA1001696	F-Y79AA1001696	0.60
5	Y79AA1001705	F-Y79AA10017Q5_	0. 67
	Y79AA1001711	F-Y79AA1001711	0.59
	Y79AA1001781	F-Y79AA1001781	0.74
40	Y79AA1001805	F-Y79AA1001805	0. 79
10	Y79AA1001827	F-Y79AA1001827	0.84
	Y79AA1001846	F-Y79AA1001846	0.34
	Y79AA1001866	F-Y79AA1001866	0. 75
15	Y79AA1001874	F-Y79AA1001874	0. 59
	Y79AA1001923	F-Y79AA1001923	0.32
	Y79AA1001963	F-Y79AA1001963	0.51
	Y79AA1002027	F-Y79AA1002027;	0.39
20	Y79AA1002083	F-Y79AA1002083	0. 65
•	Y79AA1002089	F-Y79AA1002089	0. 94
	Y79AA1002093	F-Y79AA1002093	0.89
	Y79AA1002115	F-Y79AA1002115	0.31
25	Y79AA1002125	F-Y79AA1002125	0. 76
	Y79AA1002139	F-Y79AA1002139	0. 32
	Y79AA1002204	F-Y79AA1002204	0. 94
30	Y79AA1002208	F-Y79AA1002208	0. 76
30	Y79AA1002209	F-Y79AA1002209	0. 91
	Y79AA1002210	F-Y79AA1002210	0. 60
	Y79AA1002229	F-Y79AA1002229	0. 76
35	Y79AA1002234	F-Y79AA1002234	0.94
	Y79AA1002246	F-Y79AA1002246	0.65
	Y79AA1002258	F-Y79AA1002258	0. 79
	Y79AA1002298	F-Y79AA1002298	0.83
40	Y79AA1002307	F-Y79AA1002307	0. 52
	Y79AA1002311	F-Y79AA1002311	0.92
	Y79AA1002351	F-Y79AA1002351	0.85
	Y79AA1002361	F-Y79AA1002361	0.90
45	Y79AA1002399	F-Y79AA1002399	0.94
	Y79AA1002407	F-Y79AA1002407	0.83
	Y79AA1002416	F-Y79AA1002416	0.77
50	Y79AA1002431	F-Y79AA1002431	0.61
	Y79AA1002433	F-Y79AA1002433	0.81
	Y79AA1002472	F-Y79AA1002472	0.39
	Y79AA1002482	F-Y79AA1002482	0.88
55	Y79AA1002487	F-Y79AA1002487	0. 87

Table 10

Selected clones having the maximal ATGpr1 score of not more than 0.3 (1857 clones including the 9 clones whose 5'-end sequence does not contain the ATG codon. Blank of the maximal ATGpr1 score indicates that clones whose 5'-end sequence does not contain the ATG codon).

clone	name of	maximal
name	sequence	ATGpr1
<u> </u>	···-	score
HEMBA1000	042 F-HEMBA1	000042 0.08
HEMBA1000	111 F-HEMBA1	1000111 0.1
HEMBA1000	180 F-HEMBA1	000180 0.09
HEMBA1000	213 F-HEMBA1	1000213 0.21
HEMBA1000	243 F-HEMBA1	000243 0.1
HEMBA1000	251 F-HEMBA1	1000251 0.08
HEMBA1000		
HEMBA1000	280 F-HEMBA1	1000280 0.13
HEMBA1000	282 F-HEMBA1	1000282, 0.1
HEMBA1000		
HEMBA1000		
HEMBA1000	351 F-HEMBA1	
HEMBA1000		
HEMBA1000		
HEMBA1000		
HEMBA1000	*	
HEMBA1000		1000569 0.17
HEMBA1000		

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	HEMBA1000705	F-HEMBA1000705	0.06
	HEMBA1000722	F-HEMBA1000722	0.17
	HEMBA1000726	F-HEMBA1000726	0. 08
5	HEMBA1000747	F-HEMBA1000747	0.06
	HEMBA1000749	F-HEMBA1000749	0.09
	HEMBA1000769	F-HEMBA1000769	0. 19
	HEMBA1000773	F-HEMBA1000773	0. 08
10	HEMBA1000774	F-HEMBA1000774	0. 17
,,	HEMBA1000791	F-HEMBA1000791	0. 08
	HEMBA1000737	F-HEMBA1000822	0. 19
	HEMBA1000827	F-HEMBA1000827	0. 09
45	HEMBA1000843	F-HEMBA1000827	0. 09
15	HEMBA1000852	F-HEMBA1000852	0. 12
	HEMBA1000870	F-HEMBA1000870	
	HEMBA1000876	F-HEMBA1000876	0. 15
20	HEMBA1000908	F-HEMBA1000908	0. 17
	HEMBA1000934	F-HEMBA1000934	0. 2
	HEMBA1000943	F-HEMBA1000943	0.08
	HEMBA1000960	F-HEMBA1000960	0.09
25	HEMBA1000972	F-HEMBA1000972	0. 12
	HEMBA1000974	F-HEMBA1000974	0. 08
	HEMBA1000985	F-HEMBA1000985	0.06
	HEMBA1000991	F-HEMBA1000991	0. 23
30	HEMBA1001007	F-HEMBA1001007	0. 06
	HEMBA1001017	F-HEMBA1001017	0.06
	HEMBA1001019	F-HEMBA1001019	0. 08
	HEMBA1001020	F-HEMBA1001020	0. 15
35	HEMBA1001024	F-HEMBA1001024	0. 1
	HEMBA1001026	F-HEMBA1001026	0. 13
	HEMBA1001051	F-HEMBA1001051	0. 27
	HEMBA1001060	F-HEMBA1001060	0.11
40	HEMBA1001071	F-HEMBA1001071	0. 1
	HEMBA1001077	F-HEMBA1001077	0. 07
	HEMBA1001094	F-HEMBA1001094	0. 07
	HEMBA1001099	F-HEMBA1001099	0. 08
45	HEMBA1001121	F-HEMBA1001121	0. 25
45	HEMBA1001123	F-HEMBA1001123	0. 05
	HEMBA1001172	F-HEMBA1001172	0. 08
	HEMBA1001208	F-HEMBA1001208	0.11
	HEMBA1001226	F-HEMBA1001226	0. 27
50	HEMBA1001265	F-HEMBA1001265	0.06
	HEMBA1001294	F-HEMBA1001294	0.12
	HEMBA1001299	F-HEMBA1001299	0. 2
	HEMBA1001319	F-HEMBA1001319	0. 3
55	HEMBA1001323	F-HEMBA1001323	0. 07
•	HEMBA1001327	F-HEMBA1001327	0.08

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	HEMBA1001330	F-HEMBA1001330	0.11
	HEMBA1001375	F-HEMBA1001375	0.15
	HEMBA1001383	F-HEMBA1001383	0. 11
5	HEMBA1001391	F-HEMBA1001391	0. 21
	HEMBA1001411	F-HEMBA1001411	0.06
	HEMBA1001432	F-HEMBA1001432	0. 07
	HEMBA1001433	F-HEMBA1001433	0. 24
10	HEMBA1001435	F-HEMBA1001435	0. 16
70	HEMBA1001443	F-HEMBA1001442	0. 10
	HEMBA1001454	F-HEMBA1001454	0. 11
	HEMBA1001463	F-HEMBA1001463	0. 08
15	HEMBA1001478	F-HEMBA1001478	0. 13
	HEMBA1001515		0. 3
	HEMBA1001517	,	
	HEMBA1001522	F-HEMBA1001522	0. 3
20	HEMBA1001557	F-HEMBA1001557	0.06
	HEMBA1001566	F-HEMBA1001566	0.1
	HEMBA1001585	F-HEMBA1001585	0.09
	HEMBA1001589	F-HEMBA1001589	0.06
25	HEMBA1001608	F-HEMBA1001608	0. 17
	HEMBA1001636	F-HEMBA1001636	0. 15
	HEMBA1001651	F-HEMBA1001651	0, 23
	HEMBA1001658	F-HEMBA1001658	0.17
30	HEMBA1001675	F-HEMBA1001675	0.11
30	HEMBA1001678	F-HEMBA1001678	0.06
	HEMBA1001681	F-HEMBA1001681	0.1
	HEMBA1001709	F-HEMBA1001709	0. 29
	HEMBA1001712	F-HEMBA1001712	0.08
35	HEMBA1001718	F-HEMBA1001718	0.07
	HEMBA1001734	F-HEMBA1001734	0.07
	HEMBA1001745	F-HEMBA1001745	0.09
	HEMBA1001761	F-HEMBA1001761	0.09
40	HEMBA1001784	F-HEMBA1001784	0. 11
	HEMBA1001791	F-HEMBA1001791	0. 15
,	HEMBA1001803	F-HEMBA1001803	0. 21
	HEMBA1001808	F-HEMBA1001808	0. 18
45	HEMBA1001820	F-HEMBA1001820	0.08
	HEMBA1001835	F-HEMBA1001835	0, 29
	HEMBA1001844	F-HEMBA1001844	0. 11
	HEMBA1001861	F-HEMBA1001861	0, 2
50	HEMBA1001888	F-HEMBA1001888	0. 18
	HEMBA1001918	F-HEMBA1001918	0. 10
	HEMBA1001940	F-HEMBA1001940	0. 03
	HEMBA1001942	F-HEMBA1001942	0. 17
55	HEMBA1001945	F-HEMBA1001945	0.17
55	HEMBA1001945	F-HEMBA1001945	U. UU
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	HEMBA1001962	F-HEMBA1001962	0.08
	HEMBA1001964	F-HEMBA1001964	0. 15
	HEMBA1001979	F-HEMBA1001979	0. 07
5	HEMBA1001991	F-HEMBA1001991	0.08
•	HEMBA1002003	F-HEMBA1002003	0.11
	HEMBA1002008	F-HEMBA1002008	0.08
	HEMBA1002022	F-HEMBA1002022	0.08
10	HEMBA1002039	F-HEMBA1002039	0. 2
70	HEMBA1002100	F-HEMBA1002100	0. 07
	HEMBA1002113	F-HEMBA1002113	0.09
	HEMBA1002139	F-HEMBA1002139	0. 2
45	HEMBA1002144	F-HEMBA1002144	0.11
15	HEMBA1002153	F-HEMBA1002153	0. 09
	HEMBA1002160	F-HEMBA1002160,	
	HEMBA1002162	F-HEMBA1002162	0. 19
••	HEMBA1002166	F-HEMBA1002166	0.06
20	HEMBA1002185	F-HEMBA1002185	0.11
	HEMBA1002204	F-HEMBA1002204	0.09
	HEMBA1002226	F-HEMBA1002226	0.13
•-	HEMBA1002253	F-HEMBA1002253	0. 05
25	HEMBA1002257	F-HEMBA1002257	0.09
	HEMBA1002270	F-HEMBA1002270	0.06
	HEMBA1002321	F-HEMBA1002321	0. 07
	HEMBA1002328	F-HEMBA1002328	0.06
30	HEMBA1002337	F-HEMBA1002337	0.11
	HEMBA1002348	F-HEMBA1002348	0. 12
	HEMBA1002349	F-HEMBA1002349	0. 05
	HEMBA1002381	F-HEMBA1002381	0.09
35	HEMBA1002389		0.14
	HEMBA1002486	F-HEMBA1002486	0.1
	HEMBA1002498	F-HEMBA1002498	0.17
	HEMBA1002538	F-HEMBA1002538	0. 27
40	HEMBA1002552	F-HEMBA1002552	0. 23
	HEMBA1002555	F-HEMBA1002555	0.11
	HEMBA1002558	F-HEMBA1002558	0.11
	HEMBA1002561	F-HEMBA1002561	0.11
45	HEMBA1002590	F-HEMBA1002590	0.06
	HEMBA1002592	F-HEMBA1002592	0. 17
	HEMBA1002621	F-HEMBA1002621	
	HEMBA1002628	F-HEMBA1002628	0.06
50	HEMBA1002629	F-HEMBA1002629	0. 23
	HEMBA1002645	F-HEMBA1002645 F-HEMBA1002651	0. 12 0. 08
	HEMBA1002651 HEMBA1002659	F-HEMBA1002659	0. 08
	HEMBA1002661	F-HEMBA1002661	0. 28
55		F-HEMBA1002666	0. 16
	HEMBA1002666	1 IILMDVIOUX000	U. 10

	HEMBA1002678	F-HEMBA1002678	0. 07
	HEMBA1002679	F-HEMBA1002679	0. 14
	HEMBA1002712	F-HEMBA1002712	0. 14
5	HEMBA1002716	F-HEMBA1002716	
	HEMBA1002728	F-HEMBA1002728	0. 09
	HEMBA1002730	F-HEMBA1002730	0. 03
	HEMBA1002742		
10		F-HEMBA1002742	0.09
10	HEMBA1002748 HEMBA1002780	F-HEMBA1002748	0, 17
	HEMBA1002/80	F-HEMBA1002780	0. 1 0. 1
		F-HEMBA1002801	
15	HEMBA1002826	F-HEMBA1002826	0.1
15	HEMBA1002833	F-HEMBA1002833	0. 14
	HEMBA1002886	F-HEMBA1002886	0.09
	HEMBA1002896	F-HEMBA1002896,	
	HEMBA1002921	F-HEMBA1002921	0. 25
20	HEMBA1002924	F-HEMBA1002924	0. 12
	HEMBA1002934	F-HEMBA1002934	0. 2
	HEMBA1002944	F-HEMBA1002944	0. 12
	HEMBA1002968	F-HEMBA1002968	0. 12
25	HEMBA1003034	F-HEMBA1003034	0.06
	HEMBA1003037	F-HEMBA1003037	0. 11
	HEMBA1003064	F-HEMBA1003064	0.09
	HEMBA1003071	F-HEMBA1003071	0. 27
30	HEMBA1003083	F-HEMBA1003083	0. 18
	HEMBA1003086	F-HEMBA1003086	0.06
	HEMBA1003098	F-HEMBA1003098	0.09
	HEMBA1003133	F-HEMBA1003133	0. 12
35	HEMBA1003142	F-HEMBA1003142	0. 29
	HEMBA1003166	F-HEMBA1003166	0.1
	HEMBA1003197	F-HEMBA1003197	0. 1
	HEMBA1003202	F-HEMBA1003202	0. 17
40	HEMBA1003204	F-HEMBA1003204	0. 13
	HEMBA1003212	F-HEMBA1003212	0. 13
	HEMBA1003220	F-HEMBA1003220	0. 14
	HEMBA1003229	F-HEMBA1003229	0. 19
45	HEMBA1003273	F-HEMBA1003273	0.1
40	HEMBA1003276	F-HEMBA1003276	0. 12
	HEMBA1003278	F-HEMBA1003278	0.09
	HEMBA1003296	F-HEMBA1003296	,
	HEMBA1003304	F-HEMBA1003304	0.09
50	HEMBA1003309	F-HEMBA1003309	0. 07
	HEMBA1003314	F-HEMBA1003314	0. 21
	HEMBA1003328	F-HEMBA1003328	0.13
	HEMBA1003330	F-HEMBA1003330	0.08
55	HEMBA1003348	F-HEMBA1003348	0. 17
	HEMBA1003373	F-HEMBA1003373	0. 21

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	HEMBA1003376	F-HEMBA1003376	0.14
	HEMBA1003384	F-HEMBA1003384	0.07
	HEMBA1003531	F-HEMBA1003531	0. 2
5	HEMBA1003548	F-HEMBA1003548	0.08
	HEMBA1003571	F-HEMBA1003571	0.13
	HEMBA1003579	F-HEMBA1003579	0, 13
	HEMBA1003595	F-HEMBA1003595	0.08
10	HEMBA1003597	F-HEMBA1003597	0.1
	HEMBA1003598	F-HEMBA1003598	0.12
	HEMBA1003622	F-HEMBA1003622	0.08
	HEMBA1003630	F-HEMBA1003630	0.08
15	HEMBA1003637	F-HEMBA1003637	0.07
	HEMBA1003640	F-HEMBA1003640	0.16
	HEMBA1003656	F-HEMBA1003656,	0.13
	HEMBA1003714	F-HEMBA1003714	0.12
20	HEMBA1003715	F-HEMBA1003715	0.1
	HEMBA1003725	F-HEMBA1003725	0.08
	HEMBA1003733	F-HEMBA1003733	0. 2
	HEMBA1003758	F-HEMBA1003758	0.06
25	HEMBA1003773	F-HEMBA1003773	0.11
	HEMBA1003784	F-HEMBA1003784	0.09
	HEMBA1003856	F-HEMBA1003856	0. 25
	HEMBA1003885	F-HEMBA1003885	0.09
30	HEMBA1003902	F-HEMBA1003902	0.08
•	HEMBA1003908	F-HEMBA1003908	0.08
	HEMBA1003926		0.09
	HEMBA1003937		0. 2
25	HEMBA1003939		0.12
35	HEMBA1003942		0. 1
	HEMBA1003950		0.09
	HEMBA1003958		0. 07
40	HEMBA1003976		0. 28
40	HEMBA1003987		'0.24
	HEMBA1004000		0.16
	HEMBA1004012		0.12
45	HEMBA1004015		0.23
45	HEMBA1004024		0.13
	HEMBA1004038		0.09
	HEMBA1004042		
	HEMBA1004045		0.09
50	HEMBA1004049		0.09
	HEMBA1004132		0.08
	HEMBA1004138		0.07
	HEMBA1004164		0.08
55	HEMBA1004225		0.1
	HEMBA1004241	F-HEMBA1004241	0. 08

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	HEMBA1004267	F-HEMBA1004267	0.09
	HEMBA1004272	F-HEMBA1004272	0.09
	HEMBA1004295	F-HEMBA1004295	0.19
5	HEMBA1004306	F-HEMBA1004306	0. 07
	HEMBA1004312	F-HEMBA1004312	0. 26
	HEMBA1004323	F-HEMBA1004323	0.13
	HEMBA1004334	F-HEMBA1004334	0.06
10	HEMBA1004354	F-HEMBA1004354	0.1
10	HEMBA1004356	F-HEMBA1004356	0. 07
	HEMBA1004356	F-HEMBA1004366	0.07
	HEMBA1004394	F-HEMBA1004394	0.08
15	HEMBA1004396	F-HEMBA1004396	0.05
	HEMBA1004405	F-HEMBA1004405	0.17
	HEMBA1004429	F-HEMBA1004429,	
	HEMBA1004433	F-HEMBA1004433	0. 29
20	HEMBA1004460	F-HEMBA1004460	0.11
	HEMBA1004461	F-HEMBA1004461	0.11
	HEMBA1004482	F-HEMBA1004482	0. 15
	HEMBA1004506	F-HEMBA1004506	0.05
25	HEMBA1004538	F-HEMBA1004538	0.15
25	HEMBA1004577	F-HEMBA1004577	0.17
	HEMBA1004586	F-HEMBA1004586	0.06
	HEMBA1004617	F-HEMBA1004617	0.11
	HEMBA1004629	F-HEMBA1004629	0.11
30	HEMBA1004631	F-HEMBA1004631	0.1
	HEMBA1004666	F-HEMBA1004666	0.08
	HEMBA1004670		0.15
	HEMBA1004672	F-HEMBA1004672	0.14
35	HEMBA1004730	F-HEMBA1004730	0.15
	HEMBA1004733	F-HEMBA1004733	0.12
	HEMBA1004748	F-HEMBA1004748	0.1
	HEMBA1004770	F-HEMBA1004770	0.15
40	HEMBA1004778	F-HEMBA1004778	0.14
	HEMBA1004803	F-HEMBA1004803	0.13
	HEMBA1004807	F-HEMBA1004807	0. 18
	HEMBA1004816	F-HEMBA1004816	0.06
45	HEMBA1004820	F-HEMBA1004820	0.11
40	HEMBA1004865	F-HEMBA1004865	0. 17
	HEMBA1004880	F-HEMBA1004880	
			0. 28
50	HEMBA1004900	F-HEMBA1004900	0.11
50	HEMBA1004909	F-HEMBA1004909	0.11
	HEMBA1004918	F-HEMBA1004918	0.09
	HEMBA1004956	F-HEMBA1004956	0.05
	HEMBA1004960	F-HEMBA1004960	0. 21
55	HEMBA1004978	F-HEMBA1004978	0. 1
	HEMBA1004983	F-HEMBA1004983	0. 28

	HEMBA1004995	F-HEMBA1004995	0. 24
	HEMBA1005008	F-HEMBA1005008	0. 07
	HEMBA1005039	F-HEMBA1005039	0.08
5	HEMBA1005062	F-HEMBA1005062	0.11
	HEMBA1005123	F-HEMBA1005123	0. 07
	HEMBA1005152	F-HEMBA1005152	0.11
	HEMBA1005159	F-HEMBA1005159	0.1
10	HEMBA1005232	F-HEMBA1005232	0. 1
	HEMBA1005241	F-HEMBA1005241	0. 14
	HEMBA1005251	F-HEMBA1005251	0. 29
	HEMBA1005274		0.06
15	HEMBA1005275	F-HEMBA1005275	0. 12
7.5	HEMBA1005293	F-HEMBA1005293	0. 06
	HEMBA1005304	F-HEMBA1005304,	
	HEMBA1005311	F-HEMBA1005304,	0.03
	HEMBA1005311	F-HEMBA1005311	0. 2
20	· HEMBA1005318	F-HEMBA1005318	
	HEMBA1005353		0.08
	HEMBA1005359		0. 12 0. 15
	HEMBA1005372		
25	HEMBA1005374		0.07
			0.09
	HEMBA1005389		0.1
	HEMBA1005408		0.09
30	HEMBA1005410		0. 07
	HEMBA1005426		0. 16
	HEMBA1005443		0. 17
	HEMBA1005447		0. 25
35	HEMBA1005497	F-HEMBA1005497	0.09
	HEMBA1005500	F-HEMBA1005500	0.08
	HEMBA1005506	F-HEMBA1005506	0. 17
	HEMBA1005508	F-HEMBA1005508	0. 12
40	HEMBA1005511	F-HEMBA1005511	0.1
	HEMBA1005520	F-HEMBA1005520	0.07
	HEMBA1005552	F-HEMBA1005552	0. 05
	HEMBA1005568	F-HEMBA1005568	0.06
45	HEMBA1005570	F-HEMBA1005570	0.1
45	HEMBA1005577	F-HEMBA1005577	0.13
	HEMBA1005588	F-HEMBA1005588	0. 25
	HEMBA1005593		
	HEMBA1005606	F-HEMBA1005606	0.16
50	HEMBA1005616	F-HEMBA1005616	0. 15
	HEMBA1005627	F-HEMBA1005627	0.11
		F-HEMBA1005631	0. 07
	HEMBA1005632	F-HEMBA1005632	0.07
55	HEMBA1005634	F-HEMBA1005634	0. 07
	HEMBA1005670	F-HEMBA1005670	0.11

	HEMBA1005679	F~HEMBA1005679	0. 22
	HEMBA1005699	F-HEMBA1005699	0.08
	HEMBA1005705	F-HEMBA1005705	0.08
5	HEMBA1005717	F-HEMBA1005717	0.17
	HEMBA1005765	F-HEMBA1005765	0.08
	HEMBA1005780	F-HEMBA1005780	0.16
	HEMBA1005829	F-HEMBA1005829	0. 09
10	HEMBA1005853	F-HEMBA1005853	0.06
	HEMBA1005894	F-HEMBA1005894	0.16
	HEMBA1 005909	F-HEMBA1005909	0. 13
	HEMBA1005911	F-HEMBA1005911	0. 10
15	HEMBA1005921	F-HEMBA1005921	0. 22
10	HEMBA1005931	F-HEMBA1005931	0. 16
	HEMBA1005934	F-HEMBA1005934,	
	HEMBA1005962	F-HEMBA1005962	0. 17
20	HEMBA1005999	F-HEMBA1005999	0.14
20	HEMBA1006002	F-HEMBA1006002	0. 09
	HEMBA1006005	F-HEMBA1006005	0. 2
	HEMBA1006036	F-HEMBA1006036	0. 24
05	HEMBA1006042	F-HEMBA1006042	0.11
25	HEMBA1006067	F-HEMBA1006067	0.09
	HEMBA1006090	F-HEMBA1006090	0.09
	HEMBA1006124	F-HEMBA1006124	0.05
	HEMBA1006142	F-HEMBA1006142	0.13
30	HEMBA1006158	F-HEMBA1006158	0.27
	HEMBA1006253	F-HEMBA1006253	0. 29
	HEMBA1006259	F-HEMBA1006259	0.11
	HEMBA1006268	F-HEMBA1006268	0.11
35	HEMBA1006328	F-HEMBA1006328	0.06
	HEMBA1006334	F-HEMBA1006334	0.08
	HEMBA1006359	F-HEMBA1006359	0.22
	HEMBA1006364	F-HEMBA1006364	0.23
40	HEMBA1006380	F-HEMBA1006380	0.09
	HEMBA1006416	F-HEMBA1006416	0. 25
	HEMBA1006419	F-HEMBA1006419	0.08
	HEMBA1006421	F-HEMBA1006421	0. 26
45	HEMBA1006426	F-HEMBA1006426	0.17
	HEMBA1006438	F-HEMBA1006438	0.08
	HEMBA1006446	F-HEMBA1006446	0.11
	HEMBA1006461	F-HEMBA1006461	0.15
50	HEMBA1006471	F-HEMBA1006471	0.06
	HEMBA1006486	F-HEMBA1006486	0.11
	HEMBA1006489	F-HEMBA1006489	0.09
	HEMBA1006540	F-HEMBA1006540	0. 23
55	HEMBA1006546	F-HEMBA1006546	0.07
	HEMBA1006562	F-HEMBA1006562	0.12

	HEMBA1006569	F-HEMBA1006569	
	HEMBA1006595	F-HEMBA1006595	0. 08
	HEMBA1006597	F-HEMBA1006597	0.17
5	HEMBA1006617	F-HEMBA1006617	0. 07
	HEMBA1006635	F-HEMBA1006635	0.07
	HEMBA1006639	F-HEMBA1006639	0, 25
	HEMBA1006648	F-HEMBA1006648	0. 24
10	HEMBA1006653	F-HEMBA1006653	0. 09
70	HEMBA1006665	F-HEMBA1006665	0. 03
	HEMBA1006676	F-HEMBA1006676	0. 17
	HEMBA1006695	F-HEMBA1006695	0. 17
45	HEMBA1006696	F-HEMBA1006696	0. 05
15	HEMBA1006744	F-HEMBA1006744	0. 03
	HEMBA1006779		
	HEMBA1006780	F-HEMBA1006779, F-HEMBA1006780	0. 07
	HEMBA1006760		0. 23
20		F-HEMBA1006821	
	HEMBA1006824 HEMBA1006849	F-HEMBA1006824 F-HEMBA1006849	0. 09 0. 05
	HEMBA1006865	F-HEMBA1006865	
	,,		0. 1
25	HEMBA1006921 HEMBA1006929	F-HEMBA1006921	0. 22
	HEMBA1006938	F-HEMBA1006929 F-HEMBA1006938	0. 15
	HEMBA1006938		0. 07
		F~HEMBA1006949	0. 27
30	HEMBA1007017	F-HEMBA1007017	0. 07 0. 14
	HEMBA1007045 HEMBA1007051	F-HEMBA1007045 F-HEMBA1007051	0. 14
	HEMBA1007073	F~HEMBA1007073	0. 19
	HEMBA1007078	F-HEMBA1007073	0. 22
35	HEMBA1007078	F-HEMBA1007078	0. 12
	HEMBA1007129	F-HEMBA1007129	0. 12
	HEMBA1007147	F~HEMBA1007147	0. 18
	HEMBA1007206	F-HEMBA1007206	0. 08
40	HEMBA1007256	F-HEMBA1007256	0. 11
	HEMBA1007273	F-HEMBA1007273	0. 15
	HEMBA1007279	F-HEMBA1007279	0.09
	HEMBA1007288	F-HEMBA1007288	0.06
45	HEMBA1007322	F-HEMBA1007322	0. 12
	HEMBA1007327	F-HEMBA1007327	0. 12
	HEMBA1007341	F-HEMBA1007341	0. 1
	HEMBA1007347	F-HEMBA1007347	0. 14
50	HEMBB1000005	F-HEMBB1000005	0. 13
	HEMBB1000039	F-HEMBB1000039	0. 07
	HEMBB1000044	F-HEMBB1000044	0. 08
	HEMBB1000050	F-HEMB81000050	0. 07
55	HEMBB1000054	F-HEMBB1000054	0. 12
5 5	HEMBB1000055	F-HEMB81000055	0. 2

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	HEMBB1000059	F-HEMBB1000059	0.08
	HEMBB1000089	F-HEMBB1000089	0.09
	HEMBB1000099	F-HEMBB1000099	0.17
5	HEMBB1000113	F-HEMBB1000113	0.12
	HEMBB1000141	F-HEMBB1000141	0.13
	HEMBB1000144	F-HEMBB1000144	0. 2
	HEMBB1000218	F-HEMBB1000218	0. 07
10	HEMBB1000250	F-HEMBB1000250	0. 26
70	HEMBB1000258	F-HEMBB1000258	0. 15
	HEMBB1000264	F-HEMBB1000264	0.08
	HEMBB1000272	F-HEMBB1000272	0.11
	HEMB81000274	F-HEMBB1000274	0.11
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	HEMBB1000284	F-HEMBB1000284	0.1
	HEMBB1000307	F-HEMBB1000307,	
	HEMBB1000312	F-HEMBB1000312	0.07
20	HEMBB1000317	F-HEMBB1000317	0. 27
	HEMBB1000318	F-HEMBB1000318	0.11
	HEMBB1000335	F-HEMBB1000335	0. 15
	HEMBB1000336	F-HEMBB1000336	0. 28
25	HEMBB1000337	F-HEMBB1000337	0. 25
	HEMBB1000341	F-HEMBB1000341	0. 24
	HEMBB1000343	F-HEMBB1000343	0.07
	HEMBB1000354	F-HEMBB1000354	0. 15
30	HEMBB1000369	F-HEMBB1000369	0.08
30	HEMBB1000374	F-HEMB81000374	0. 3
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	HEMBB1000399	F-HEMBB1000399	0.08
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35	HEMBB1000404	F-HEMBB1000404	0. 29
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	HEMBB1000434	F-HEMBB1000434	0.05
	HEMBB1000441	F-HEMBB1000441	0.12
40	HEMB81000455	F-HEMBB1000455	0.05
•	HEMBB1000472	F-HEMBB1000472	0.05
	HEMBB1000480	F-HEMBB1000480	0.13
	HEMBB1000487	F-HEMB81000487	0.09
45	HEMBB1000490	F-HEMBB1000490	0.06
	HEMBB1000491	F-HEMBB1000491	0.16
	HEMBB1000493	F-HEMBB1000493	0.15
	HEMBB1000518	F-HEMBB1000518	0.05
50	HEMBB1000523	F-HEMBB1000523	0.07
	HEMBB1000530	F-HEMBB1000530	0.14
	HEMBB1000554	F-HEMBB1000554	0. 23
	HEMBB1000564	F-HEMBB1000564	0.09
	HEMBB1000573	F-HEMBB1000573	0. 03
55	HEMBB1000575	F-HEMBB1000575	0. 08
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	HEMBB1000598	F-HEMBB1000598	0.12
	HEMBB1000637	F-HEMBB1000637	0.19
5	HEMBB1000638	F-HEMB81000638	0.07
	HEMBB1000643	F-HEMBB1000643	0.07
	HEMBB1000649	F-HEMBB1000649	0.13
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10	HEMBB1000665	F-HEMBB1000665	0.07
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	HEMBB1000726	F-HEMBB1000726	0. 07
45	HEMBB1000738	F-HEMBB1000738	0. 19
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	HEMBB1000790	F-HEMBB1000790	0. 23
	HEMBB1000794	F-HEMB81000794	0.09
20	HEMBB1000821	F-HEMBB1000821	0.03
	HEMBB1000821	F-HEMBB1000822	0.18
	HEMBB1000827	F-HEMBB1000827	0.10
	HEMBB1000831	F-HEMBB1000831	0. 1
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	HEMBB1000870	F-HEMB81000883	0. 12
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	HEMBB1000893	F-HEMB81000893	0. 03
	HEMBB1000910	F-HEMBB1000910	0. 06
	HEMBB1000913	F-HEMBB1000913	0.09
35	HEMBB1000915	F-HEMBB1000915	0.12
	HEMBB1000917	F-HEMBB1000917	0. 15
	HEMBB1000959	F-HEMBB1000959	0.08
	HEMBB1000981	F-HEMBB1000981	0.09
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	HEMBB1001008	F-HEMBB1001008	0. 25
	HEMBB1001037	F-HEMBB1001037	0. 14
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	HEMB81001119	F-HEMBB1001119	0.13
	HEMBB1001113	F-HEMBB1001113	0.13
	HEMBB1001142	F-HEMB81001142	0.17
55	HEMBB1001177	F-HEMBB1001177	0. 17
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	HEMBB1001209	F-HEMBB1001209	0.11
	HEMBB1001218	F-HEMBB1001218	0.1
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	HEMBB1001249	F-HEMBB1001249	0.12
	HEMBB1001253	F-HEMB81001253	0.13
	HEMBB1001254	F-HEMBB1001254	0.13
10	HEMBB1001267	F-HEMBB1001267	0.09
	HEMBB1001271	F-HEMB81001271	0, 16
	HEMBB1001282	F-HEMB81001282	0.14
	HEMBB1001302	F-HEMBB1001302	0.18
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-	HEMBB1001315	F-HEMBB1001315	0. 18
	HEMBB1001317	F-HEMBB1001317,	
	HEMBB1001326	F-HEMBB1001326	0. 09
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	HEMBB1001586	F-HEMBB1001586	0.06
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•	HEMB81002617	F-HEMBB1002617	0. 12
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	HEMBB1002635	F-HEMBB1002635	0.08
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	HEMBB1002683	F-HEMB81002683	0.06
	HEMBB1002686	F-HEMBB1002686	0. 12
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	HEMBB1002702	F-HEMBB1002702	0.16
	HEMBB1002712	F-HEMBB1002712	0. 07
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	MAMMA1000092	F-MAMMA1000092	0.11
	MAMMA1000103	F-MAMMA1000103	0.09
55	MAMMA1000117	F-MAMMA1000117	0. 2
	MAMMA1000129	F-MAMMA1000129	0. 17

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	MAMMA1000133	F-MAMMA1000133	0.1
	MAMMA1000134	F-MAMMA1000134	0.05
	MAMMA1000155	F-MAMMA1000155	0.12
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v	MAMMA1000198	F-MAMMA1000198	0. 2
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15	MAMMA1000270	F-MAMMA1000270	0.1
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	MAMMA1000356	F-MAMMA1000356	0. 15
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50	MAMMA1000597		0.1
	MAMMA1000605	F-MANNA1000605	0.09
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E 6	MAMMA1000643	F-MAMMA1000643	0. 1
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	MAMMA1000696	F-MAMMA1000696	0.08
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10	MAMMA1000733	F-MAMMA1000733	0.1
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20	MAMMA1000839	F-MAMMA1000839	0. 11
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	MAMMA1000855		0. 08
25	MAMMA1000862	F-MAMMA1000855 F-MAMMA1000862	0. 23
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			0. 16
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30	MAMMA1000876	F-MAMMA1000876 F-MAMMA1000877	0.14
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	MAMMA1000940	F-MAMMA1000940	0. 21
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40	MAMMA1000943	F-MAMMA1000943	0.09
	MAMMA1000957	F-MAMMA1000957	0.13
	MAMMA1000962	F-MAMMA1000962	0.06
	MAMMA1000975	F-MAMMA1000975	0. 26
45	MAMMA1000987	F-MAMMA1000987	0.08
	MAMMA1000998	F-MAMMA1000998	0.08
	MAMMA1001003	F-MAMMA1001003	0. 15
	MAMMA1001024	F-MAMMA1001024	0. 08
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	MAMMA1001035	F-MAMMA1001035	0.08
	MAMMA1001038	F-MAMMA1001038	0. 15
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55	MAMMA1001067	F-MAMMA1001067	0.08
	MAMMA1001074	F-MAMMA1001074	0. 09

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	MAMMA1001082 F-MAMMA1001082	0. 1
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20	MAMMA1001330 F-MAMMA1001330	0. 13
	MAMMA1001341 F-MAMMA1001341	0. 16
	MAMMA1001383 F-MAMMA1001383	0. 05
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	MAMMA1001575 F-MAMMA1001575	0.08
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	MAMMA1001606 F-MAMMA1001606	0.13
45	MANNA1001663 F-MANNA1001663	0.11
	MANNA1001670 F-MANNA1001670	0.2
	MAMMAT001671 F-MAMMA1001671	0.13
	MAMMA1001679 F-MAMMA1001679	0.13
50	MAMMA1001683 F-MAMMA1001683	0.16
30	MAMMA1001686 F-MAMMA1001686	0. 09
	MAMMA1001711 F-MAMMA1001711	0. 11
	MANNA1001715 F-MANNA1001715	0.11
	MAMMA1001740 F-MAMMA1001740	0. 1
55	MAMMA1001745 F-MAMMA1001745	0.09
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5	MAMMA1001806	F-MANIMA1001806	0.06
	MAMMA1001815	F-MAMMA1001815	0.1
	MAMMA1001818	F-MAMMA1001818	0.1
	MAMMA1001820	F-MAMMA1001820	0. 27
10	MAMMA1001836	F-MAMMA1001836	0.05
	MAMMA1001854	F-MAMMA1001854	0.17
	MAMMA1001878	F-MAMMA1001878	0, 13
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19	MAMMA1001907	F-MAMMA1001907	0. 00
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	MAMMA1001931	F-MAMMA1001931	0. 3
	MAMMA1001951	F-MAMMA1001963	0. 14
20	MAMMA1001993	F-MAMMA1001992	0. 08
			0. 12
	MAMMA1002032	F-MAMMA1002032	
	MAMMA1002056	F-MAMMA1002056	0.09
25	MAMMA1002058	F-MAMMA1002058	0.16
	MAMMA1002078	F-MAMMA1002078	0. 28
	MAMMA1002082	F-MAMMA1002082	0.08
	MAMMA1002084	F-MAMMA1002084	0. 24
30		F-MAMMA1002093	0. 12
		F-MAMMA1002108	0.05
		F-MAMMA1002118	0. 1
	MAMMA1002125	F-MAMMA1002125	0.09
35	MAMMA1002132	F-MAMMA1002132	0.11
	MAMMA1002140	F-MAMMA1002140	0.11
	MAMMA1002145	F-MAMMA1002145	0.11
	MAMMA1002155	F-MAMMA1002155	0.1
40	MAMMA1002158	F-MAMMA1002158	0.11
40	MAMMA1002215	F-MAMMA1002215	0.05
	MAMMA1002230	F-MAMMA1002230	0.07
	MAMMA1002250	F-MAMMA1002250	0. 2
	MAMMA1002267	F-MAMMA1002267	0.08
45	MAMMA1002282	F-MAMMA1002282	0. 05
	MAMMA1002293	F-MAMMA1002293	0.09
	MAMMA1002298	F-MAMMA1002298	0.09
	MAMMA1002299	F-MAMMA1002299	0. 16
50	MAMMA1002310	F-MAMMA1002310	0. 16
	MAMMA1002311	F-MAMMA1002311	0. 13
	MAMMA1002322	F-MAMMA1002322	0. 07
	MAMMA1002332	F-MAMMA1002332	0.08
55	MAMMA1002339	F-MANMA1002339	0.09
	MAMMA1002347	F-MAMMA1002347	0.06

	MAMMA1002352	F-MAMMA1002352	0. 08
	MAMMA1002359	F-MAMMA1002359	0. 08
	MAMMA1002360	F-MAMMA1002360	0. 13
5	MAMMA1002361		
•		F-MAMMA1002361	
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	MAMMA1002411		0. 28
10	MAMMA1002417	F-MAMMA1002417	0. 19
10	MAMMA1002428	F-MAMMA1002428	0.07
	MAMMA1002446	F-MAMMA1002446	0. 17
	MAMMA1002475	F-MAMMA1002475	0. 17
45	MAMMA1002480	F-MAMMA1002480	0. 07
15	MAMMA1002494	F-MAMMA1002494	0.17
	MAMMA1002498	F-MAMMA1002498	0. 28
	MAMMA1002545	F-MAMMA1002545	
	MAMMA1002556	F-MAMMA1002556	0.06
20	MAMMA1002566	F-MAMMA1002566	0. 13
	MAMMA1002571	F-MAMMA1002571	0. 29
	MAMMA1002573	F-MAMMA1002573	0. 29
	MAMMA1002597	F-MAMMA1002597	0.1
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	MAMMA1002612	F-MAMMA1002612	0.14
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	MAMMA1002622	F-MAMMA1002622	0. 24
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	MAMMA1002629	F-MAMMA1002629	0. 05
	MAMMA1002636	F-MAMMA1002636	0. 12
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	MAMMA1002698	F-MAMMA1002698	0.08
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	MAMMA1002727	F-MAMMA1002727	0. 15
	MAMMA1002728	F-MAMMA1002728	0.12
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50	MAMMA1002758	F-MAMMA1002758	0. 29
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	MAMMA1002765	F-MAMMA1002765	0. 29
	MAMMA1002780	F-MAMMA1002780	0.12
	MAMMA1002820	F-MAMMA1002820	0.3
55	MAMMA1002830	F-MAMMA1002830	0.16
	MAMMA1002833	F-MAMMA1002B33	0.14

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	MAMMA1002835	F-MAMMA1002835	0. 13
	MAMMA1002838	F-MAMMA1002838	0.09
	MAMMA1002844	F-MAMMA1002844	0. 27
5	MAMMA1002858	F-MAMMA1002858	0. 29
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	MAMMA1002880	F-MAMMA1002880	0.11
	MAMMA1002887	F-MAMMA1002887	0.09
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	MAMMA1002908	F-MAMMA1002908	0.11
	MAMMA1002909	F-MAMMA1002909	0. 22
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25	MAMMA1003019	F-MAMMA1003019	0. 12
	MAMMA1003026	F-MAMMA1003026	0. 26
	MAMMA1003031	F-MAMMA1003031	0.18
	MAMMA1003039	F-MAMMA1003039	0.09
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	MAMMA1003089	F-MAMMA1003089	0.14
	MAMMA1003140	F-MAMMA1003140	0.19
	NT2RM2000609	F-NT2RM2000609	0. 26
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	NT2RM4000086	F-NT2RM4000086	0. 28
	NT2RM4000200	F-NT2RM4000200	0.16
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	NT2RM4000265	F-NT2RM4000265	0.18
	NT2RM4000327	F-NT2RM4000327	0.06
	NT2RM4000366	F-NT2RM4000366	0. 28
45	NT2RM4000368	F-NT2RM4000368	0.11
	NT2RM4000414	F-NT2RM4000414	0.19
	NT2RM4000425	F-NT2RM4000425	0. 22
	NT2RM4000532	F-NT2RM4000532	0.14
	NT2RM4000779	F-NT2RM4000779	0.13
50	NT2RM4000855	F-NT2RM4000855	0.13
	NT2RM4000979	F-NT2RM4000979	0. 22
	NT2RM4001016	F-NT2RM4001016	0. 2
	NT2RM4001160	F-NT2RM4001160	0. 1
55	NT2RM4001313	F-NT2RM4001313	0. 26

	NT2RM4001414	F-NT2RM4001414	0. 23
	NT2RM4001437	F-NT2RM4001437	0. 12
	NT2RM4001519	F-NT2RM4001519	0. 07
5	NT2RM4001557	F-NT2RM4001557	
	NT2RM4001605	F-NT2RM4001605	0. 22
		F-NT2RM4001754	0. 22
	NT2RM4001754		
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	NT2RM4002287	F-NT2RM4002287	0. 27
	NT2RM4002383	F-NT2RM4002383	0. 08
	NT2RM4002390	F-NT2RM4002390	0. 18
25	NT2RM4002438	F-NT2RM4002438	0. 23
	NT2RM4002479	F-NT2RM4002479	0. 27
	NT2RM4002499	F-NT2RM4002499	0.08
	NT2RM4002504	F-NT2RM4002504	0.08
20	NT2RM4002532	F-NT2RM4002532	0. 15
30	NT2RM4002567	F-NT2RM4002567	0. 28
	NT2RP2000027	F-NT2RP2000027	0.07
	NT2RP2000076	F-NT2RP2000076	0.16
	NT2RP2000077	F-NT2RP2000077	0.2
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	NT2RP2000108	F-NT2RP2000108	0.07
	NT2RP2000183	F-NT2RP2000183	0.12
	NT2RP2000257	F-NT2RP2000257	0. 27
40	NT2RP2000258	F-NT2RP2000258	0.19
	NT2RP2000270	F-NT2RP2000270	0.06
	NT2RP2000289	F-NT2RP2000289	0.06
	NT2RP2000327	F-NT2RP2000327	0.08
45	NT2RP2000337	F-NT2RP2000337	0.14
	NT2RP2000420	F-NT2RP2000420	0. 2
	NT2RP2000459	F-NT2RP2000459	0.12
	NT2RP2000498	F-NT2RP2000498	0.05
50	NT2RP2000523	F-NT2RP2000523	0.14
50	NT2RP2000603	F-NT2RP2000603	0. 22
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	NT2RP2000678	F-NT2RP2000678	0.11
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5	NT2RP2000987	F-NT2RP2000987	0. 12
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	NT2RP2001196	F-NT2RP2001196	0.08
10	NT2RP2001226	F-NT2RP2001226	0. 18
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	NT2RP2001295	F-NT2RP2001295	0. 13
	NT2RP2001312	F-NT2RP2001312	0. 15
15	NT2RP2001347	F-NT2RP2001347	0.11
	NT2RP2001423	F-NT2RP2001423	0.06
	NT2RP2001445	F-NT2RP2001445	
	NT2RP2001449	F-NT2RP2001449	0. 15
20	NT2RP2001467	F-NT2RP2001467	0. 21
	NT2RP2001506	F-NT2RP2001506	0.06
	NT2RP2001526	F-NT2RP2001526	0. 1
	NT2RP2001569	F-NT2RP2001569	0. 09
25	NT2RP2001663	F-NT2RP2001663	0.08
	NT2RP2001677	F-NT2RP2001677	0. 14
	NT2RP2001678	F-NT2RP2001678	0.17
	NT2RP2001699	F-NT2RP2001699	0. 05
30	NT2RP2001720	F-NT2RP2001720	0.08
	NT2RP2001740	F-NT2RP2001740	0. 23
	NT2RP2001762	F-NT2RP2001762	0.11
	NT2RP2001813	F-NT2RP2001813	0. 19
35	NT2RP2001869	F-NT2RP2001869	0. 22
	NT2RP2001926	F-NT2RP2001926	0. 07
	NT2RP2001936	F-NT2RP2001936	0.08
	NT2RP2001943	F-NT2RP2001943	0. 27
40	NT2RP2002032	F-NT2RP2002032	0. 23
40	NT2RP2002041	F-NT2RP2002041	0. 1
	NT2RP2002047	F-NT2RP2002047	0. 12
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45	NT2RP2002105	F-NT2RP2002105	0.13
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	NT2RP2002172	F-NT2RP2002172	0. 21
50	NT2RP2002192	F-NT2RP2002192	0.05
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	NT2RP2002256	F-NT2RP2002256	0.17
	NT2RP2002259	F-NT2RP2002259	0. 28
55	NT2RP2002316	F-NT2RP2002316	0. 15
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	NT2RP2002439	F-NT2RP2002439	0.07
	NT2RP2002457	F-NT2RP2002457	0.11
	NT2RP2002475	F-NT2RP2002475	0.14
5	NT2RP2002504	F-NT2RP2002504	0, 21
	NT2RP2002546	F-NT2RP2002546	0.17
	NT2RP2002591	F-NT2RP2002591	0. 15
	NT2RP2002606	F-NT2RP2002606	0. 11
40	NT2RP2002643	F-NT2RP2002643	0. 26
10	NT2RP2002043	F-NT2RP2002727	0. 26
		F-NT2RP2002727	0. 20
	NT2RP2002736		
	NT2RP2002740	F-NT2RP2002740	0.16
15	NT2RP2002741	F-NT2RP2002741	0.11
	NT2RP2002750	F-NT2RP2002750	0.14
	NT2RP2002752	F-NT2RP2002752	
	NT2RP2002753	F-NT2RP2002753	0.09
20	NT2RP2002778	F-NT2RP2002778	0.13
	NT2RP2002839	F-NT2RP2002839	
	NT2RP2002857	F-NT2RP2002857	0. 21
	NT2RP2002987	F-NT2RP2002987	0.08
25	NT2RP2003000	F-NT2RP2003000	0. 24
25	NT2RP2003073	F-NT2RP2003073	0.08
	NT2RP2003129	F-NT2RP2003129	0. 07
	NT2RP2003161	F-NT2RP2003161	0.05
	NT2RP2003164	F-NT2RP2003164	0.09
30	NT2RP2003206	F-NT2RP2003206	0. 22
	NT2RP2003230	F-NT2RP2003230	0.08
	NT2RP2003237	F-NT2RP2003237	0. 24
	NT2RP2003339	F-NT2RP2003339	0.1
35	NT2RP2003394	F-NT2RP2003394	0.07
	NT2RP2003456	F-NT2RP2003456	0.08
	NT2RP2003499	F-NT2RP2003499	0.08
	NT2RP2003517	F-NT2RP2003517	0. 29
40	NT2RP2003522	F-NT2RP2003522	0. 22
	NT2RP2003559	F-NT2RP2003559	0.12
	NT2RP2003668	F-NT2RP2003668	0.08
	NT2RP2003704	F-NT2RP2003704	0.09
45	NT2RP2003706	F-NT2RP2003706	0. 28
45	NT2RP2003727	F-NT2RP2003727	0. 17
	NT2RP2003770	F-NT2RP2003770	0.09
	NT2RP2003859	F-NT2RP2003859	0. 18
	NT2RP2003871	F-NT2RP2003871	0. 23
50	NT2RP2003885	F-NT2RP2003885	0. 24
	NT2RP2003912	F-NT2RP2003912	0, 26
	NT2RP2003968	F-NT2RP2003968	0, 2
	NT2RP2003988	F-NT2RP2003988	0.09
55	NT2RP2004014	F-NT2RP2004014	0.07

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	NT2RP2004170	F-NT2RP2004170	0.09
5	NT2RP2004172	F-NT2RP2004172	0.15
	NT2RP2004207	F-NT2RP2004207_	0.11
	NT2RP2004232	F-NT2RP2004232	0.17
	NT2RP2004270	F-NT2RP2004270	0. 29
	NT2RP2004300	F-NT2RP2004300	0.11
10	NT2RP2004321	F-NT2RP2004321	0.13
	NT2RP2004339	F-NT2RP2004339	0.14
	NT2RP2004347	F-NT2RP2004347	0. 14
	NT2RP2004396	F-NT2RP2004396	0. 29
15	NT2RP2004400	F-NT2RP2004400	0.06
	NT2RP2004412	F-NT2RP2004412	0. 24
	NT2RP2004425	F-NT2RP2004425	0. 13
	NT2RP2004490	F-NT2RP2004490	0. 13
20	NT2RP2004512	F-NT2RP2004512	0. 1
	NT2RP2004523	F-NT2RP2004523	0. 15
	NT2RP2004580	F-NT2RP2004580	0. 12
	NT2RP2004587	F-NT2RP2004587	0. 08
25	NT2RP2004675	F-NT2RP2004675	0.14
20	NT2RP2004681	F-NT2RP2004681	0. 17
	NT2RP2004709	F-NT2RP2004709	0.08
	NT2RP2004736	F-NT2RP2004736	0.13
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30	NT2RP2004775	F-NT2RP2004775	0.17
	NT2RP2004961	F-NT2RP2004961	0. 21
	NT2RP2004962	F-NT2RP2004962	0. 24
	NT2RP2004967	F-NT2RP2004967	0.05
35	NT2RP2004982	F-NT2RP2004982	0.11
	NT2RP2005020	F-NT2RP2005020	0.16
	NT2RP2005031	F-NT2RP2005031	0.07
	NT2RP2005108	F-NT2RP2005108	0.11
40	NT2RP2005254	F-NT2RP2005254	0.16
	NT2RP2005289	F-NT2RP2005289	0. 22
	NT2RP2005293	F-NT2RP2005293	0.08
	NT2RP2005325	F-NT2RP2005325	
45	NT2RP2005336	F-NT2RP2005336	0. 27
	NT2RP2005344	F-NT2RP2005344	0. 24
	NT2RP2005354	F-NT2RP2005354	0.08
	NT2RP2005360	F-NT2RP2005360	0.12
50	NT2RP2005407	F-NT2RP2005407	0. 28
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	NT2RP2005491	F-NT2RP2005491	0.18
	NT2RP2005501	F-NT2RP2005501	0. 1
55	NT2RP2005531	F-NT2RP2005531	0. 29
55	NT2RP2005581	F-NT2RP2005581	0.07

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	NT2RP2005645	F-NT2RP2005645	0.13
	NT2RP2005651	F-NT2RP2005651	0.06
	NT2RP2005694	F-NT2RP2005694	0. 3
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	NT2RP2005857	F-NT2RP2005857	0. 28
	NT2RP2005908	F-NT2RP2005908	0. 05
15	NT2RP2005942	F-NT2RP2005942	0. 26
	NT2RP2005980	F-NT2RP2005980,	
	NT2RP2006098	F-NT2RP2006098	0.06
	NT2RP2006103	F-NT2RP2006103	0.07
20	NT2RP2006166	F-NT2RP2006166	0. 27
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	NT2RP2006261	F-NT2RP2006261	0. 16
25	NT2RP2006320	F-NT2RP2006320	0. 16
	NT2RP2006321	F~NT2RP2006321	0. 15
	NT2RP2006323	F-NT2RP2006323	0. 10
	NT2RP2006393	F-NT2RP2006393	0. 07
30	NT2RP2006454	F-NT2RP2006454	0. 11
	NT2RP2006467	F-NT2RP2006467	0. 24
	NT2RP2006534	F-NT2RP2006534	0. 14
	NT2RP2006554	F-NT2RP2006554	0.16
35	NT2RP2006598	F-NT2RP2006598	0. 08
	NT2RP3000055	F-NT2RP3000055	0. 14
	NT2RP3000109	F-NT2RP3000109	0. 18
	NT2RP3000186	F-NT2RP3000186	0.06
40	NT2RP3000233	F-NT2RP3000233	0. 29
40 .	NT2RP3000235	F-NT2RP3000235	0. 09
	NT2RP3000247	F-NT2RP3000247	0. 19
	NT2RP3000341	F-NT2RP3000341	0. 16
	NT2RP3000418	F-NT2RP3000418	0. 15
45	NT2RP3000433	F-NT2RP3000433	0. 11
	NT2RP3000441	F-NT2RP3000441	0. 2
	NT2RP3000449	F-NT2RP3000449	0.16
	NT2RP3000451	F-NT2RP3000451	0.14
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	NT2RP3000542	F-NT2RP3000542	0. 1
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55	NT2RP3000582	F-NT2RP3000582	0.08

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	NT2RP3000622	F-NT2RP3000622	0.17
	NT2RP3000628	F-NT2RP3000628	0.08
5	NT2RP3000685	F-NT2RP3000685	0.17
	NT2RP3000736	F-NT2RP3000736	0. 26
	NT2RP3000742	F-NT2RP3000742	0. 22
	NT2RP3000815	F-NT2RP3000815	0.06
10	NT2RP3000865	F-NT2RP3000865	0.13
	NT2RP3000875	F-NT2RP3000875	0.16
	NT2RP3000904	F-NT2RP3000904	0.13
	NT2RP3001007	F-NT2RP3001007	0.1
15	NT2RP3001115	F-NT2RP3001115	0.1
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	NT2RP3001281	F-NT2RP3001281	0. 29
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20	NT2RP3001339	F-NT2RP3001339	0. 27
	NT2RP3001340	F-NT2RP3001340	0.11
	NT2RP3001374	F-NT2RP3001374	0. 21
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25	NT2RP3001432	F-NT2RP3001432	0. 29
	NT2RP3001459	F-NT2RP3001459	0.16
	NT2RP3001527	F-NT2RP3001527	0.3
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30	NT2RP3001589	F-NT2RP3001589	0.12
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10	PLACE1008630 F-PLACE1008630 0.12
	PLACE1008643 F-PLACE1008643 0.11
	PLACE1008715 F-PLACE1008715 0.2
	PLACE1008748 F-PLACE1008748 0.15
15	PLACE1008757 F-PLACE1008757 0.13
	PLACE1008798 F-PLACE1008798, 0.14
	PLACE1008807 F-PLACE1008807 0.12
	PLACE1008851 F-PLACE1008851 0.22
20	PLACE1008920 F-PLACE1008920 0.1
	PLACE1008941 F-PLACE1008941 0.14
	PLACE1008947 F-PLACE1008947 0.05
	PLACE1009020 F-PLACE1009020 0.14
25	PLACE1009039 F-PLACE1009039 0.16
	PLACE1009048 F-PLACE1009048 0.09
	PLACE1009050 F-PLACE1009050 0.09
	PLACE1009150 F-PLACE1009150 0.12
30	PLACE1009155 F-PLACE1009155 0.08
	PLACE1009172 F-PLACE1009172 0.07
	PLACE1009183 F-PLACE1009183 0.07
	PLACE 1009200 FT LAGE 1000
35	PLACE 1009246
	PLACE 1009300 F F LAGE 100000
	PLACE 1009390 1 1 CROC 100000
	PLACE 1009410 T TENDETOS
40	PLAGE 1009477 T TEMBER OF CO.
	PLACETO09493
	PLACE 1009009 1 1 2 10 2 10 11
	PLACE 1009333
45	PLACE 1009613 1 1 LACE 1000621 0 19
	1 0 07
	121021000000000000000000000000000000000
	PLACE1009639 F-PLACE1009639 0.27 PLACE1009879 F-PLACE1009879 0.07
50	PLACE1009879 F-PLACE1009888 0.06
	PLACE1009924 F-PLACE1009924 0.09
	PLACE1009925 F-PLACE1009925 0.21
	PLACE1009935 F-PLACE1009935 0.22
55	PLACE1009947 F-PLACE1009947 0.19
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	PLACE1010069	F-PLACE1010069	0. 22
	PLACE1010083	F-PLACE1010083	0. 26
	PLACE1010089	F-PLACE1010089	0, 13
5	PLACE1010181	F-PLACE1010181	0. 07
	PLACE1010231	F-PLACE1010231	0.17
	PLACE1010270	F-PLACE1010270	0.08
	PLACE1010341	F-PLACE1010341	0.1
10	PLACE1010562	F-PLACE1010562	0, 14
10	PLACE1010579	F-PLACE1010579	0.15
	PLACE1010624	F-PLACE1010624	0.13
	PLACE1010628	F-PLACE1010628	0.16
	PLACE1010631	F-PLACE1010631	0. 10
15	PLACE1010631	F-PLACE1010662	0. 23
	PLACE1010702	F-PLACE1010702,	
	PLACE1010739	F-PLACE1010739	0. 13
20	PLACE1010802	F-PLACE1010802	0. 17
	PLACE1010891	F-PLACE1010891	0.09
	PLACE1010896	F-PLACE1010896	0. 22
	PLACE1010916	F-PLACE1010916	0. 15
25	PLACE1010947	F-PLACE1010947	0. 08
	PLACE1011032	F-PLACE1011032	0. 17
	PLACE1011057	F-PLACE1011057	0. 1
	PLACE1011109	F-PLACE1011109	0. 08
30	PLACE1011133	F-PLACE1011133	0. 08
30	PLACE1011143	F-PLACE1011143	0. 23
	PLACE1011165	F-PLACE1011165	0. 05
	PLACE1011185	F-PLACE1011185	0. 17
	PLACE1011203	F-PLACE1011203	0.08
35	PLACE1011296	F-PLACE1011296	0.13
	PLACE1011340	F-PLACE1011340	0. 05
	PLACE1011375	F-PLACE1011375	0.07
	PLACE1011419	F-PLACE1011419	0.15
40	PLACE1011452	F-PLACE1011452	0.13
	PLACE1011465	F-PLACE1011465	0.14
	PLACE1011492	F-PLACE1011492	0. 26
	PLACE1011503	F-PLACE1011503	0.14
45	PLACE1011520	F-PLACE1011520	0.12
	PLACE1011567	F-PLACE1011567	0. 13
	PLACE1011576	F-PLACE1011576	0. 29
	PLACE1011641	F-PLACE1011641	0.07
50	PLACE1011643	F-PLACE1011643	0. 27
	PLACE1011649	F-PLACE1011649	0.17
	PLACE1011650	F-PLACE1011650	0.13
	PLACE1011719	F-PLACE1011719	0.15
55	PLACE1011729	F-PLACE1011729	0. 23
	PLACE1011749	F-PLACE1011749	0.12

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	PLACE1011762	F-PLACE1011762	0. 06
	PLACE1011778	F-PLACE1011778	0. 07
	PLACE1011874	F-PLACE1011874	0. 07
5	PLACE1011962	F-PLACE1011962	0.06
	PLACE1011964	F-PLACE1011964	0. 13
	PLACE1011995	F-PLACE1011995	0. 08
	PLACE2000003	F-PLACE2000003	0. 09
10	PLACE2000011	F-PLACE2000011	0. 12
10	PLACE2000017	F-PLACE2000017	0. 22
	PLACE2000061	F-PLACE2000061	0. 19
	PLACE2000103	F-PLACE2000103	0.16
	PLACE2000115	F-PLACE2000115	0.11
15	PLACE2000113	F-PLACE2000132	0. 71
	PLACE2000132	F-PLACE2000136,	
	PLACE2000130	F-PLACE2000140	0. 24
	PLACE2000140		0. 24
20	PLACE2000176	F-PLACE2000176	0. 25
		F-PLACE2000187	
	PLACE2000216 PLACE2000264	F-PLACE2000216	0. 15
		F-PLACE2000264	0.06
25	PLACE2000305	F-PLACE2000305	0.05
	PLACE2000317	F-PLACE2000317	0. 25
	PLACE2000335	F-PLACE2000335	0.09
	PLACE2000342	F-PLACE2000342	0.07
30	PLACE2000347	F-PLACE2000347	0. 12
	PLACE2000366	F-PLACE2000366	0. 08
	PLACE2000379		0. 24
	PLACE2000394	F-PLACE2000394	0. 2
35	PLACE2000398	F-PLACE2000398	0. 21
35	PLACE2000419		0. 05
	PLACE2000425	F-PLACE2000425	0. 16
	PLACE2000435	F-PLACE2000435	0. 05
	PLACE2000450	F-PLACE2000450	0. 12
40	PLACE2000455	F-PLACE2000455	0. 15
	PLACE2000465	F-PLACE2000465	0. 12
	PLACE2000477	F-PLACE2000477	0. 3
	PLACE3000004	F-PLACE3000004	0. 3
45	PLACE3000119	F-PLACE3000119	0. 1
	PLACE3000124	F-PLACE3000124	0.11
	PLACE3000136	F-PLACE3000136	0. 13
	PLACE3000158	F-PLACE3000158	0. 06
50	PLACE3000169	F-PLACE3000169	0.13
	PLACE3000199	F-PLACE3000199	0. 06
	PLACE3000207	F-PLACE3000207	0. 11
	PLACE3000208	F-PLACE3000208	0. 07
55	PLACE3000218	F-PLACE3000218	0. 15
	PLACE3000220	F-PLACE3000220	0. 05

	PLACE3000230	F-PLACE3000230	0. 23
	PLACE3000271	F-PLACE3000271	0.17
	PLACE3000276	F-PLACE3000276	0. 2
5	PLACE3000341	F-PLACE3000341	0.08
	PLACE3000362	F-PLACE3000362	0. 19
	PLACE3000365	F-PLACE3000365	0. 09
	PLACE3000373	F-PLACE3000373	0. 22
10	PLACE3000388	F-PLACE3000388	0. 12
,,	PLACE3000399	F-PLACE3000399	0. 14
	PLACE3000400	F-PLACE3000400	0. 08
	PLACE3000401	F-PLACE3000401	0. 1
	PLACE3000401	F-PLACE3000401	0. 1
15		F-PLACE3000402	0. 1
	PLACE3000406		
	PLACE3000475	F-PLACE3000475,	
	PLACE4000093	F-PLACE4000093	0. 21
20	PLACE4000100	F-PLACE4000100	0. 19
		F-PLACE4000233	0.07
	PLACE4000247	F-PLACE4000247	0. 14
	PLACE4000250	F-PLACE4000250	0. 17
25	PLACE4000252	F-PLACE4000252	0.09
	PLACE4000320	F-PLACE4000320	0. 23
	PLACE4000344	F-PLACE4000344	0. 21
	PLACE4000367	F-PLACE4000367	0. 18
30	PLACE4000379	F-PLACE4000379	0. 14
30	PLACE4000401	F-PLACE4000401	0. 13
	PLACE4000411	F-PLACE4000411	0. 11
	PLACE4000494	F-PLACE4000494	0.19
	PLACE4000548	F-PLACE4000548	0. 27
35	THYR01000026	F-THYR01000026	0.09
	THYR01000035	F-THYR01000035	0.09
	THYR01000085	F-THYR01000085	0.13
	THYR01000092	F-THYR01000092	0. 21
40	THYR01000111	F-THYR01000111	0. 13
·	THYR01000129	F-THYR01000129	0.11
	THYR01000132	F-THYR01000132	0. 1
	THYR01000156	F-THYR01000156	0. 13
45	THYR01000186	F-THYR01000186	0. 12
	THYR01000187	F-THYR01000187	0.11
	THYR01000190	F-THYR01000190	0.16
	THYR01000221	F-THYR01000221	0.08
50	THYR01000241	F-THYR01000241	0.08
	THYR01000279	F-THYR01000279	0. 13
	THYR01000438	F-THYR01000438	0.06
	THYR01000452		0. 12
	THYR01000471	F-THYR01000471	0. 11
55	THYR01000484	F-THYR01000484	0. 24
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	THYR01000558	F-THYR01000558	0.09
	THYR01000596	F-THYR01000596	0.14
	THYR01000602	F-THYR01000602	0.13
5	THYR01000625	F-THYR01000625	0. 2
	THYR01000641	F-THYR01000641	0.06
	THYR01000658	F-THYR01000658	0. 12
	THYR01000699	F-THYR01000699	0.09
10	THYR01000748	F-THYR01000748	0. 3
10	THYR01000793	F-THYR01000793	0. 25
	THYR01000796	F-THYR01000796	0. 23
	THYR01000790	F-THYR01000805	0. 11
	THYR01000805	F-THYR01000805	0. 14
15			0.14
	THYR01000843	F-THYR01000843	
	THYR01000852	F-THYR01000852,	
	THYR01000865	F-THYR01000865	0.14
20	THYR01000895	F-THYR01000895	0.12
	THYR01000974	F-THYR01000974	0. 07
	THYR01000975	F-THYR01000975	0.08
	THYR01001003	F-THYR01001003	0.11
25	THYR01001031	F-THYR01001031	0.1
	THYR01001062	F-THYR01001062	0. 14
	THYR01001093	F-THYR01001093	0.13
	THYR01001121	F-THYR01001121	0.14
30	THYR01001133	F-THYR01001133	0. 24
	THYR01001177	F-THYR01001177	
	THYR01001262	F-THYR01001262	0.2
	THYR01001290	F-THYR01001290	0.06
	THYR01001321	F-THYR01001321	0.09
35	THYR01001322	F-THYR01001322	0.13
	THYR01001365	F-THYR01001365	0.08
	THYR01001401	F-THYR01001401	0.21
	THYR01001411	F-THYR01001411	0.06
40	THYR01001426	F-THYR01001426	0. 1
	THYR01001434	F-THYR01001434	0. 15
	THYR01001480	F-THYR01001480	0. 1
	THYR01001534	F-THYR01001534	0. 26
45	THYR01001541	F-THYR01001541	0. 22
	THYR01001559	F-THYR01001559	0.15
	THYR01001570	F-THYR01001570	0, 28
	THYR01001573	F-THYR01001573	0. 05
50	THYR01001595	F-THYR01001595	0. 05
	THYR01001617	F-THYR01001617	0. 1
	THYR01001673	F-THYR01001673	0. 07
	THYR01001706	F-THYR01001706	0. 15
ee.	THYR01001745	F-THYR01001745	0. 12
55	THYR01001746	F-THYR01001746	0. 1

	THYR01001772	F-THYR01001772	0.08
	THYR01001895	F-THYR01001895	0. 19
	THYR01001907	F-THYR01001907	0. 17
	Y79AA1000033	F-Y79AA1000033	0.18
	Y79AA1000065	F-Y79AA1000065	0. 22
	Y79AA1000346	F-Y79AA1000346	0. 2
	Y79AA1000405	F-Y79AA1000405	0.07
	Y79AA1000410	F-Y79AA1000410	0.08
	Y79AA1000538	F-Y79AA1000538	0.06
	Y79AA1000802	F-Y79AA1000802	0. 28
	Y79AA1000805	F-Y79AA1000805	0. 2
	Y79AA1000969	F-Y79AA1000969	0. 07
	Y79AA1001061	F-Y79AA1001061	0. 22
	Y79AA1001167	F-Y79AA1001167,	0. 22
	Y79AA1001384	F-Y79AA1001384	0. 18
•	Y79AA1001594	F-Y79AA1001594	0. 22
	Y79AA1001692	F-Y79AA1001692	0. 2
	Y79AA1001848	F-Y79AA1001848	0. 21
	Y79AA1001875	F-Y79AA1001875	0.09
	Y79AA1002103	F-Y79AA1002103	0. 11
	Y79AA1002211	F-Y79AA1002211	0. 19
	Y79AA1002220	F-Y79AA1002220	0. 25

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Table 11

574 clones belonging to groups (4), (5), (6), and (7), each having the maximal ATGpr1 score of not more than 0.3 and selected based on the maximal ATGpr2 score, and their maximal ATGpr2 scores

clone	name of	maximal
name	sequence	ATGpr2
		score
HEMBA100018	0 F-HEMBA1000180	0. 37
HEMBA100025	1 F-HEMBA1000251	0. 14
HEMBA100028	0 F-HEMBA1000280	0. 33
HEMBA100028	2 F-HEMBA1000282	0. 36
HEMBA100033	3 F-HEMBA1000333	0.59
HEMBA100035	1 F-HEMBA1000351	0.36
HEMBA100035	7 F-HEMBA1000357	0.39
HEMBA100037	6 F-HEMBA1000376	0. 53
HEMBA100046	9 F-HEMBA1000469	0.32
HEMBA100050	4 F-HEMBA1000504	0. 34
HEMBA100051	9 F-HEMBA1000519	0. 72
HEMBA100054	5 F-HEMBA1000545	0.32
HEMBA100055	7 F-HEMBA1000557	0. 35
HEMBA100057	5 F-HEMBA1000575	0.34

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	HEMBA1000604	F-HEMBA1000604	0. 28
	HEMBA1000622	F-HEMBA1000622	0.32
	HEMBA1000673	F-HEMBA1000673	0.14
5	HEMBA1000726	F-HEMBA1000726	0.32
	HEMBA1000749	F-HEMBA1000749	0. 35
	HEMBA1000822	F-HEMBA1000822	0, 42
	HEMBA1000876	F-HEMBA1000876	0. 44
10	HEMBA1000934	F-HEMBA1000934	0. 41
	HEMBA1000943	F-HEMBA1000943	0. 34
	HEMBA1000960	F-HEMBA1000960	0.34
	HEMBA1000972	F-HEMBA1000972	0. 44
15	HEMBA1001020	F-HEMBA1001020	0. 31
	HEMBA1001024	F-HEMBA1001024	0. 22
	HEMBA1001026	F-HEMBA1001026,	0.28
••	HEMBA1001051	F-HEMBA1001051	0.36
20	HEMBA1001060	F-HEMBA1001060	0.34
	HEMBA1001071	F-HEMBA1001071	0.57
	HEMBA1001208	F-HEMBA1001208	0.57
	HEMBA1001226	F-HEMBA1001226	0. 44
25	HEMBA1001319	F-HEMBA1001319	0.58
	HEMBA1001383	F-HEMBA1001383	0. 35
	HEMBA1001391	F-HEMBA1001391	0. 42
	HEMBA1001411	F-HEMBA1001411	0. 55
30	HEMBA1001433	F-HEMBA1001433	0. 66
	HEMBA1001435	F-HEMBA1001435	0.72
	HEMBA1001478	F-HEMBA1001478	0. 36
	HEMBA1001522	F-HEMBA1001522	0.31
35	HEMBA1001636	F-HEMBA1001636	0.32
	HEMBA1001651	F-HEMBA1001651	0. 54
	HEMBA1001658	F-HEMBA1001658	0. 38
	HEMBA1001709	F-HEMBA1001709	0. 55
40	HEMBA1001734	F-HEMBA1001734	0. 13
	HEMBA1001745	F-HEMBA1001745	0. 33
	HEMBA1001784	F-HEMBA1001784	0.08
	HEMBA1001791	F-HEMBA1001791	0.48
45	HEMBA1001803	F-HEMBA1001803	0. 21
,,,	HEMBA1001808	F-HEMBA1001808	0. 23
	HEMBA1001835	F-HEMBA1001835	0. 67
	HEMBA1001844	F-HEMBA1001844	0. 46
50	HEMBA1001888	F-HEMBA1001888	0. 72
50	HEMBA1001918	F-HEMBA1001918	0. 25
	HEMBA1001940	F-HEMBA1001940	0. 40
	HEMBA1001962	F-HEMBA1001962	0.63
	HEMBA1002022	F-HEMBA1002022	0. 32
55	HEMBA1002185	F-HEMBA1002185	0.33
	HEMBA1002257	F-HEMBA1002257	0.13
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	HEMBA1002348	F-HEMBA1002348	0.39
	HEMBA1002621	F-HEMBA1002621	0. 47
	HEMBA1002645	F-HEMBA1002645	0.34
5	HEMBA1002661	F-HEMBA1002661	
	HEMBA1002728	F-HEMBA1002728	
	HEMBA1002742	F-HEMBA1002742	0. 49
	HEMBA1002921	F-HEMBA1002921	0.79
10	HEMBA1002924	F-HEMBA1002924	0.45
	HEMBA1002934	F-HEMBA1002934	
	HEMBA1002968	F-HEMBA1002968	0.39
	HEMBA1003037	F-HEMBA1003037	0.43
15	HEMBA1003064	F-HEMBA1003064	0. 19
	HEMBA1003071		0.48
	HEMBA1003166	F-HEMBA1003166,	
	HEMBA1003202	F-HEMBA1003202	0.38
20	HEMBA1003204	F-HEMBA1003204	0.34
20	HEMBA1003220	F-HEMBA1003220	0.39
	HEMBA1003229	F-HEMBA1003229	0, 43
	HEMBA1003276	F-HEMBA1003276	0,48
25	HEMBA1003296	F-HEMBA1003296	0.36
	HEMBA1003373	F-HEMBA1003373	0.48
	HEMBA1003531	F-HEMBA1003531	0.46
	HEMBA1003640	F-HEMBA1003640	0.32
30	HEMBA1003714	F-HEMBA1003714	0.16
30	HEMBA1003856	F-HEMBA1003856	0.47
	HEMBA1003902	F-HEMBA1003902	0.17
	HEMBA1003908	F-HEMBA1003908	0.44
35	HEMBA1003926	F-HEMBA1003926	0.36
33	HEMBA1003942		0. 47
	HEMBA1003987	F-HEMBA1003987	0. 39
	HEMBA1004000		0. 87
40	HEMBA1004024		0.82
40	HEMBA1004164	F-HEMBA1004164	0. 21
	HEMBA1004267	F-HEMBA1004267	0. 46
	HEMBA1004323		0. 37
	HEMBA1004405	F-HEMBA1004405	0. 23
45	HEMBA1004433		0.51
•	HEMBA1004577	F-HEMBA1004577	0. 51
	HEMBA1004730	F-HEMBA1004730	0.55
	HEMBA1004778	F-HEMBA1004778	0. 47
50	HEMBA1004803	F-HEMBA1004803	0.36
	HEMBA1004807	F-HEMBA1004807	0.48
	HEMBA1004880	F-HEMBA1004880	0.43
	HEMBA1004900	F-HEMBA1004900	0.40
55	HEMBA1004909	F-HEMBA1004909	0. 24
	HEMBA1004983	F-HEMBA1004983	0. 35

	HEMBA1005123	F-HEMBA1005123	0. 35
	HEMBA1005232	F-HEMBA1005232	0. 09
	HEMBA1005241	F-HEMBA1005241	0. 34
5	HEMBA1005311	F-HEMBA1005311	
	HEMBA1005318	F-HEMBA1005318	0. 59
	HEMBA1005353	F-HEMBA1005353	0. 34
	HEMBA1005374	F-HEMBA1005374	0. 38
10	HEMBA1005447	F-HEMBA1005447	0. 34
70	HEMBA1005508	F-HEMBA1005508	0. 14
	HEMBA1005577	F-HEMBA1005577	0. 14
			0. 22
	HEMBA1005588	F-HEMBA1005588	
15	HEMBA1005593	F-HEMBA1005593	0. 44
	HEMBA1005606	F-HEMBA1005606	0. 38
	HEMBA1005679	F-HEMBA1005679,	
	HEMBA1005894	F-HEMBA1005894	
20	HEMBA1005911	F-HEMBA1005911	0. 56
	HEMBA1006036	F-HEMBA1006036	0. 87
	HEMBA1006124	F-HEMBA1006124	0. 35
	HEMBA1006253	F-HEMBA1006253	0. 35
25	HEMBA1006259	F-HEMBA1006259	0. 54
	HEMBA1006364	F-HEMBA1006364	0. 55
	HEMBA1006380	F-HEMBA1006380	0. 48
	HEMBA1006426	F-HEMBA1006426	0. 54
30	HEMBA1006461	F-HEMBA1006461	0. 20
	HEMBA1006562	F-HEMBA1006562	0. 72
	HEMBA1006597	F-HEMBA1006597	0.64
	HEMBA1006639	F-HEMBA1006639	0. 55
or.	HEMBA1006653	F-HEMBA1006653	0. 34
35	HEMBA1006695	F-HEMBA1006695	0. 16
	HEMBA1006696	F-HEMBA1006696	0. 37
	HEMBA1006744	F-HEMBA1006744	0. 33
	HEMBA1006824	F-HEMBA1006824	0. 37
40	HEMBA1006949	F-HEMBA1006949	0. 78
	HEMBA1007078	F-HEMBA1007078	0. 44
	HEMBA1007129	F-HEMBA1007129	0. 55
	HEMBA1007147	F-HEMBA1007147	0. 41
45	HEMBA1007206	F-HEMBA1007206	0. 46
	HEMBA1007279	F-HEMBA1007279	0. 45
	HEMBA1007327	F-HEMBA1007327	0.71
	HEMBB1000005	F-HEMBB1000005	0. 35
50	HEMBB1000054	F-HEMBB1000054	0.18
	HEMBB1000055	F-HEMBB1000055	0. 87
	HEMBB1000144	F-HEMB81000144	0. 47
	HEMBB1000258	F-HEMBB1000258	0. 87
55	HEMBB1000318	F-HEMBB1000318	0. 34
	HEMBB1000335	F-HEMBB1000335	0. 48

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	HEMBB1000336	F-HEMBB1000336	0. 35
	HEMBB1000354	F-HEMBB1000354	0. 47
	HEMBB1000374	F-HEMBB1000374	0. 69
5	HEMBB1000402	F-HEMBB1000402	0. 54
	HEMBB1000404	F-HEMBB1000404	0.47
	HEMBB1000480	F-HEMBB1000480	0. 35
	HEMBB1000493	F-HEMBB1000493	0. 33
10	HEMBB1000554	F-HEMBB1000554	0.46
	HEMBB1000573	F-HEMBB1000573	0.34
	HEMBB1000649	F-HEMBB1000649	0. 41
	HEMBB1000652	F-HEMBB1000652	0. 46
15	HEMBB1000709	F-HEMBB1000709	0. 56
,,	HEMBB1000749	F-HEMBB1000749	0. 36
	HEMBB1000790	F-HEMBB1000790,	0. 58
	HEMBB1000827	F-HEMBB1000827	0. 31
20	HEMBB1000831	F-HEMBB1000831	0.42
20	HEMBB1000893	F-HEMBB1000893	0. 36
	HEMBB1001004	F-HEMBB1001004	0. 43
	HEMBB1001008	F-HEMBB1001008	0. 44
. =	HEMBB1001047	F-HEMBB1001047	0. 47
25	HEMBB1001119	F-HEMBB1001119	0. 25
	HEMBB1001142	F-HEMBB1001142	0.15
	HEMBB1001315	F-HEMBB1001315	0. 72
	HEMBB1001317	F-HEMBB1001317	0. 49
30	HEMBB1001326	F-HEMBB1001326	0. 44
	HEMBB1001337	F-HEMBB1001337	0.18
	HEMBB1001367	F-HEMBB1001367	0. 35
	HEMBB1001424	F-HEMBB1001424	0. 34
35	HEMBB1001436	F-HEMBB1001436	0.34
	HEMBB1001458	F-HEMBB1001458	0. 47
	HEMBB1001535	F-HEMBB1001535	0.34
	HEMB81001536	F-HEMB81001536	0. 22
40	HEMBB1001565	F-HEMBB1001565	0. 35
	HEMBB1001747	F-HEMBB1001747	0.40
	HEMBB1001749	F-HEMBB1001749	0. 39
	HEMBB1001797	F-HEMBB1001797	0. 79
45	HEMBB1001836	F-HEMBB1001836	0. 52
	HEMBB1001863	F-HEMBB1001863	0. 38
	HEMB81001868	F-HEMBB1001868	0. 27
	HEMBB1001875	F-HEMBB1001875	0. 46
50	HEMBB1001911	F-HEMBB1001911	0.87
	HEMBB1001922	F-HEMBB1001922	0. 38
	HEMBB1001925	F-HEMBB1001925	0. 33
	HEMBB1001944	F-HEMB81001944	0. 35
55	HEMBB1001983	F-HEMB81001983	0.32
	HEMBB1001996	F-HEMBB1001996	0.39

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	HEMBB1001997	F-HEMBB1001997	0.36
	HEMBB1002045	F-HEMBB1002045	0. 20
	HEMBB1002092	F-HEMBB1002092	0. 31
5	HEMBB1002247	F-HEMB81002247	0.53
·	HEMBB1002266	F-HEMBB1002266	0. 34
	HEMBB1002387	F-HEMBB1002387	0. 32
	HEMBB1002425	F-HEMBB1002425	0. 31
	HEMBB1002423	F-HEMBB1002458	0. 44
10			
	HEMBB1002522	F-HEMBB1002522	0. 56
	HEMBB1002534	F-HEMBB1002534	0. 43
	HEMBB1002579	F-HEMBB1002579	0. 21
15	HEMBB1002582	F-HEMBB1002582	0, 48
	HEMBB1002596	F-HEMBB1002596	0. 42
	HEMBB1002617	F-HEMBB1002617,	0. 47
	HEMBB1002702	F-HEMBB1002702	0.31
20	MAMMA1000043	F-MAMMA1000043	0.45
20	MAMMA1000092	F-MAMMA1000092	0.39
	MAMMA1000129	F-MAMMA1000129	0. 47
	MAMMA1000175	F-MAMMA1000175	0. 18
	MAMMA1000198	F-MAMMA1000198	0. 46
25	MAMMA1000221	F-MAMMA1000221	0. 43
	MAMMA1000307	F-MAMMA1000307	0. 35
	MAMMA1000331	F-MAMMA1000331	0. 41
	MAMMA1000356	F-MAMMA1000356	0. 41
30	MAMMA1000360	F-MAMMA1000360	0. 33
	MAMMA1000402	F-MAMMA1000402	0. 35
	MAMMA1000414	F-MAMMA1000414	0. 32
	MAMMA1000444	F-MAMMA1000444	0. 13
35	MAMMA1000500	F-MAMMA1000500	0. 36
	MAMMA1000522	F-MAMMA1000522	0.46
	MAMMA1000576	F-MAMMA1000576	0. 58
	MAMMA1000594	F-MAMMA1000594	0.69
40	MAMMA1000597	F-MAMMA1000597	0. 35
40	MAMMA1000720	F-MAMMA1000720	0. 43
	MAMMA1000720	F-MAMMA1000720	0. 43
		F-MAMMA1000761	
	MAMMA1000775	F-MAMMA1000778	0. 35 0. 54
45	MAMMA1000778		
	MAMMA1000798	F-MAMMA1000798	0.40
	MAMMA1000862	F-MAMMA1000862	0. 87
	MAMMA1000876	F-MAMMA1000876	0. 47
50	MAMMA1000883	F-MAMMA1000883	0.41
	MAMMA1000931	F-MAMMA1000931	0. 45
	MAMMA1000940	F-MAMMA1000940	0. 35
	.MAMMA1000941	F-MAMMA1000941	0. 35
55	MAMMA1000943	F-MAMMA1000943	0. 23
	MAMMA1000975	F-MAMMA1000975	0. 32

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	MAMMA1001038	F-MAMMA1001038	0.41
	MAMMA1001162	F-MAMMA1001162	0. 18
	MAMMA1001186	F-MAMMA1001186	0.49
5	MAMMA1001220		
	MAMMA1001256		0. 35
	MAMMA1001274		0. 57
	MAMMA1001341		0. 45
-	MAMMA1001341		
10		F-MAMMA1001397	0. 64
	MAMMA1001420		0. 41
	MAMMA1001547		0. 39
	MAMMA1001590	F-MAMMA1001590	0. 46
15	MAMMA1001670	F-MAMMA1001670	0. 36
	MAMMA1001679	F-MAMMA1001679	0.46
	MAMMA1001711	F-MAMMA1001711,	0.68
	MAMMA1001745	F-MAMMA1001745	0.53
20	MAMMA1001760	F-MAMMA1001760	0.58
20	MAMMA1001769	F-MAMMA1001769	0.33
	MAMMA1001815	F-MAMMA1001815	0.43
	MAMMA1001820	F-MAMMA1001820	0.19
	MAMMA1001907	F-MAMMA1001907	0. 38
25	MAMMA1002056	F-MAMMA1002056	0.35
	MAMMA1002078		0. 34
	MAMMA1002093		0. 55
	MAMMA1002125		0.39
30	MAMMA1002132		0. 57
	MAMMA1002145		0.78
	MAMMA1002250		0.72
	MAMMA1002311	F-MAMMA1002311	0.58
35	MAMMA1002360	F-MAMMA1002360	0. 19
	MAMMA1002411	F-MAMMA1002411	0. 40
	MAMMA1002498	F-MAMMA1002498	0. 37
	MAMMA1002438	F-MAMMA1002571	0. 37
40	MAMMA1002771	F-MAMMA1002701	0. 55
70	MAMMA1002707	F-MAMMA1002707	0. 34
	MAMMA1002728	F-MAMMA1002728	0. 55
	MAMMA1002746		0. 45
45	MAMMA1002764		0. 44
	MAMMA1002765	F-MAMMA1002765	0. 40
	MAMMA1002820	F-MAMMA1002820	0. 32
	MAMMA1002830	F-MAMMA1002830	0. 36
50	MAMMA1002909	F-MAMMA1002909	0. 64
	MAMMA1002941	F-MAMMA1002941	0. 42
	MAMMA1002972	F-MAMMA1002972	0.08
	MAMMA1002973	F-MAMMA1002973	0. 35
55	MAMMA1003004	F-MAMMA1003004	0. 42
	MAMMA1003007	F-MAMMA1003007	0. 41

	MAMMA1003039	F-MAMMA1003039	0.46
	MAMMA1003089	F-MAMMA1003089	0. 58
	NT2RM4000086	F-NT2RM4000086	0.36
5	NT2RM4000265	F-NT2RM4000265	
	NT2RM4000366	F-NT2RM4000366	0. 21
•	NT2RM4000414	F-NT2RM4000414	0. 35
	NT2RM4000425	F-NT2RM4000425	0. 22
10	NT2RM4000779	F-NT2RM4000779	0. 43
	NT2RM4000855	F-NT2RM4000855	0. 52
	NT2RM4001160	F-NT2RM4001160	0. 36
	NT2RM4001313	F-NT2RM4001313	0. 38
15	NT2RM4001437	F-NT2RM4001437	0. 53
	NT2RM4001754	F-NT2RM4001754	0. 35
	NT2RM4001856	F-NT2RM4001856,	
	NT2RM4001953	F-NT2RM4001953	0, 59
20	NT2RM4001984	F-NT2RM4001984	0. 35
	NT2RM4002390	F-NT2RM4002390	0. 24
	NT2RP2000077	F-NT2RP2000077	0. 35
	NT2RP2000108	F-NT2RP2000108	0. 08
25	NT2RP2000183	F-NT2RP2000183	0. 43
	NT2RP2000257	F-NT2RP2000257	0. 07
	NT2RP2000289	F-NT2RP2000289	0. 35
	NT2RP2000420	F-NT2RP2000420	0. 48
30	NT2RP2000678	F-NT2RP2000678	0. 55
	NT2RP2000715	F-NT2RP2000715	0. 37
	NT2RP2000842	F-NT2RP2000842	0.40
	NT2RP2000970	F-NT2RP2000970	0. 46
35	NT2RP2001149	F-NT2RP2001149	0.31
	NT2RP2001226	F-NT2RP2001226	0. 44
	NT2RP2001295	F-NT2RP2001295	0.40
	NT2RP2001347	F-NT2RP2001347	0. 41
40	NT2RP2001467	F-NT2RP2001467	0. 44
	NT2RP2001506	F-NT2RP2001506	0.17
	NT2RP2001569	F-NT2RP2001569	0. 31
·	NT2RP2001663	F-NT2RP2001663	0. 37
45	NT2RP2001936	F-NT2RP2001936	0. 33
	NT2RP2002041	F-NT2RP2002041	0. 31
	NT2RP2002047	F-NT2RP2002047	0. 15
	NT2RP2002066	F-NT2RP2002066	0.11
50	NT2RP2002172	F-NT2RP2002172	0. 32
	NT2RP2002219	F-NT2RP2002219	0. 54
	NT2RP2002316	F-NT2RP2002316	0.39
	NT2RP2002475	F-NT2RP2002475	0.09
55	NT2RP2002546	F-NT2RP2002546	0. 33
	NT2RP2002591	F-NT2RP2002591	0.38
	NT2RP2002643	F-NT2RP2002643	0. 55

		NT2RP2002741	F-NT2RP2002741	0.73
		NT2RP2002750	F-NT2RP2002750	0.48
		NT2RP2002778	F-NT2RP2002778	0.41
5		NT2RP2002857	F-NT2RP2002857_	0.34
		NT2RP2003000	F-NT2RP2003000	0.56
		NT2RP2003073	F-NT2RP2003073	0.36
		NT2RP2003237	F-NT2RP2003237	0.69
1	0	NT2RP2003394	F-NT2RP2003394	0.34
		NT2RP2003517	F-NT2RP2003517	0.44
		NT2RP2003668	F-NT2RP2003668	0.46
		NT2RP2003988	F-NT2RP2003988	0. 55
1	5	NT2RP2004232	F-NT2RP2004232	0, 34
		NT2RP2004400	F-NT2RP2004400	0.18
		NT2RP2004512	F-NT2RP2004512,	
		NT2RP2004523	F-NT2RP2004523	0. 72
2	0	NT2RP2004587	F-NT2RP2004587	0.09
_		NT2RP2004736	F-NT2RP2004736	0. 70
		NT2RP2004767	F-NT2RP2004767	0. 54
		NT2RP2004775	F-NT2RP2004775	0. 53
2	5	NT2RP2004961	F-NT2RP2004961	0.49
~		NT2RP2004962	F-NT2RP2004962	0. 87
		NT2RP2004982	F-NT2RP2004982	0. 32
		NT2RP2005289	F-NT2RP2005289	0. 24
3	0	NT2RP2005407	F-NT2RP2005407	0. 33
3		NT2RP2005531	F-NT2RP2005531	0. 20
		NT2RP2005694	F-NT2RP2005694	0. 21
		NT2RP2005726	F-NT2RP2005726	0. 42
3	s.	NT2RP2005942	F-NT2RP2005942	0.10
3	3	NT2RP2006258	F-NT2RP2006258	0.37
		NT2RP2006261	F-NT2RP2006261	0.42
		NT2RP2006454	F-NT2RP2006454	0.46
4	•	NT2RP2006554	F-NT2RP2006554	0.14
4		NT2RP3000055	F-NT2RP3000055	0.31
		NT2RP3000233	F-NT2RP3000233	0.54
		NT2RP3000341	F-NT2RP3000341	0.43
	-	NT2RP3000418	F-NT2RP3000418	0.53
4	5	NT2RP3000451	F-NT2RP3000451	0.46
		NT2RP3000561	F-NT2RP3000561	0.47
		NT2RP3000582	F-NT2RP3000582	0.44
		NT2RP3001007	F-NT2RP3001007	0.26
5	U	NT2RP3001281	F-NT2RP3001281	0.36
		NT2RP3001318	F-NT2RP3001318	0. 21
		NT2RP3001339	F-NT2RP3001339	0.58
		NT2RP3001340	F-NT2RP3001340	0. 41
5	5	NT2RP3001383	F-NT2RP3001383	0. 36
		NT2RP3001432	F-NT2RP3001432	0.34

	NT2RP3001580	F-NT2RP3001580	0. 45
	NT2RP3001589	F-NT2RP3001589	0.48
	NT2RP3001898	F-NT2RP3001898	0.19
5	NT2RP3002004	F-NT2RP3002004	0.36
	NT2RP3002173	F-NT2RP3002173	0. 39
	NT2RP3003133	F-NT2RP3003133	0. 35
	NT2RP3003264	F-NT2RP3003264	0.13
10	NT2RP3003346	F-NT2RP3003346	0.72
•	NT2RP3003403	F-NT2RP3003403	0. 45
	NT2RP3003433	F-NT2RP3003433	0. 24
	NT2RP3003576	F-NT2RP3003576	0. 35
15	NT2RP3003625	F-NT2RP3003625	0. 43
7.5	NT2RP3003665	F-NT2RP3003665	0. 32
	NT2RP3003800	F-NT2RP3003800,	
	NT2RP3003828	F-NT2RP3003828	0.52
00	NT2RP3003842	F-NT2RP3003842	0. 32
20	NT2RP3004070	F-NT2RP3004070	0. 23
	NT2RP3004070	F-NT2RP3004470	0. 46
	NT2RP4000023	F-NT2RP4000023	0. 33
	NT2RP4000025	F-NT2RP4000025	0. 56
25	NT2RP4000102	F-NT2RP4000102	0. 38
	NT2RP4000167	F-NT2RP4000167	0. 42
	NT2RP4000167	F-NT2RP4000167	0. 42
	NT2RP4000214	F-NT2RP4000214	0. 33
30	NT2RP4000424	F-NT2RP4000424	0.37
	NT2RP4000424	F-NT2RP4000915	0. 46
	NT2RP4000915	F-NT2RP4000915	0. 37
	OVARC1000017	F-0VARC1000017	0.43
35	OVARC1000017	F-0VARC1000017	0.14
	OVARC1000085	F-0VARC1000085	0. 17
	OVARC1000085	F-0VARC1000085	0.58
	OVARC1000092	F-0VARC1000092	0.53
40	OVARC1000145	F-0VARC1000145	0. 53
	OVARC1000414	F-0VARC1000414	0.40
	OVARC1000486	F-0VARC1000486	0. 15
	0VARC1000496	F-0VARC1000496	0. 33
45	0VARC1000526	F-0VARC1000948	0. 33
	OVARC1000948	F-0VARC1000948	0. 71
	OVARC1001240		
50	OVARC1001600 OVARC1001805	F-0VARC1001600	0.37
		F-0VARC1001805	0.51
	OVARC1001813	F-0VARC1001813	0. 58
	OVARC1001846	F-0VARC1001846	0.59
55	PLACE1000540	F-PLACE1000540	0.35
	PLACE1000599	F-PLACE1000599	0.37
	PLACE1001088	F-PLACE1001088	0. 36

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	PLACE1001323	F-PLACE1001323	0. 21
	PLACE1001377	F-PLACE1001377	0. 41
	PLACE1001440	F-PLACE1001440	0. 45
5	PLACE1001517	F-PLACE1001517	0.41
	PLACE1001672	F-PLACE1001672	0. 35
	PLACE1001705	F-PLACE1001705	0.11
	PLACE1001756	F-PLACE1001756	0.67
10	PLACE1002157	F-PLACE1002157	0.68
	PLACE1002205	F-PLACE1002205	0.55
	PLACE1002227	F-PLACE1002227	0.14
	PLACE1002259	F-PLACE1002259	0.47
45	PLACE1002319	F-PLACE1002319	0. 29
15	PLACE1002319	F-PLACE1002319	0.23
	PLACE1002399	F-PLACE1002477,	
	PLACE1002477	F-PLACE1002477	0. 35
	PLACE1002583		
20		F-PLACE1002583	0.49
	PLACE1002604	F-PLACE1002604	0. 25
	PLACE1002772	F-PLACE1002772	0. 13
	PLACE1002853	F-PLACE1002853	0. 46
25	PLACE1002968	F-PLACE1002968	0. 33
•	PLACE1003238	F-PLACE1003238	0. 46
	PLACE1003420	F-PLACE1003420	0. 42
	PLACE1003478	F-PLACE1003478	0. 20
30	PLACE1003566	F-PLACE1003566	0.35
	PLACE1003593	F-PLACE1003593	0. 42
	PLACE1003618	F-PLACE1003618	0.36
	PLACE1004274	F-PLACE1004274	0.41
35	PLACE1004681	F-PLACE1004681	0. 25
	PLACE1004716	F-PLACE1004716	0.49
	PLACE1004773	F-PLACE1004773	0.31
	PLACE1004815	F-PLACE1004815	0.72
40	PLACE1004836	F-PLACE1004836	0.39
40	PLACE1004913	F-PLACE1004913	0.44
	PLACE1004979	F-PLACE1004979	0.44
	PLACE1005052	F-PLACE1005052	0. 63
45	PLACE1005086	F-PLACE1005086	0. 57
45	PLACE1005108	F-PLACE1005108	0.17
	PLACE1005128	F-PLACE1005128	0.40
	PLACE1005176	F-PLACE1005176	0. 48
	PLACE1005467	F-PLACE1005467	0. 37
50	PLACE1005639	F-PLACE1005639	0. 33
	PLACE1005850	F-PLACE1005850	0. 54
	PLACE1005932	F-PLACE1005932	0. 20
	PLACE1006003	F-PLACE1006003	0. 71
55	PLACE1006017	F-PLACE1006017	0. 42
,	PLACE1006288	F-PLACE1006288	0.34

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	PLACE1006318	F-PLACE1006318	0. 22
	PLACE1006368	F-PLACE1006368	0.10
	PLACE1006371	F-PLACE1006371	0. 43
5	PLACE1006506	F-PLACE1006506	
	PLACE1006629	F-PLACE1006629	0.44
	PLACE1006795	F-PLACE1006795	0.35
•	PLACE1006904	F-PLACE1006904	0, 20
10	PLACE1007478	F-PLACE1007478	0. 70
	PLACE1007557	F-PLACE1007557	0. 17
	PLACE1007743	F-PLACE1007743	0. 28
	PLACE1007829		0. 16
15	PLACE1007877	F-PLACE1007877	0. 16
13	PLACE1008181	F-PLACE1008181	0. 39
	PLACE1008330	F-PLACE1008330,	
	PLACE1008584	F-PLACE1008584	0. 61
00	PLACE1008630	F-PLACE1008630	0.09
20	PLACE1008715	F-PLACE1008715	0. 36
	PLACE1008851	F-PLACE1008851	0. 39
	PLACE1008941	F-PLACE1008941	0. 42
	PLACE1009039	F-PLACE1009039	0. 43
25	PLACE1009048	F-PLACE1009048	0. 24
	PLACE1009493	F-PLACE1009493	0.35
	PLACE1009539		0.31
	PLACE1009637	F-PLACE1009637	0. 58
30	PLACE1009925	F-PLACE1009925	0.17
	PLACE1009947	F-PLACE1009947	0. 42
	PLACE1010231	F-PLACE1010231	0. 36
	PLACE1010562	F-PLACE1010562	0. 63
35	PLACE1010579	F-PLACE1010579	0. 42
	PLACE1010739	F-PLACE1010739	0.49
	PLACE1010802	F-PLACE1010802	0.41
	PLACE1010896	F-PLACE1010896	0. 47
40	PLACE1011032	F-PLACE1011032	0.46
	PLACE1011109	F-PLACE1011109	0. 19
	PLACE1011185	F-PLACE1011185	0.66
	PLACE1011452	F-PLACE1011452	0. 72
45	PLACE1011465	F-PLACE1011465	0. 51
	PLACE1011492	F-PLACE1011492	0.19
	PLACE1011520	F-PLACE1011520	0. 54
	PLACE1011567	F-PLACE1011567	0.51
50	PLACE1011643	F-PLACE1011643	0. 21
	PLACE10117.19	F-PLACE1011719	0. 55
	PLACE1011749	F-PLACE1011749	0. 24
	PLACE2000011	F-PLACE2000011	0. 48
55	PLACE2000017	F-PLACE2000017	0. 34
	PLACE2000061	F-PLACE2000061	0. 34

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10		, , =). 47
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	PLACE2000477	, , _,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	0.36
	PLACE3000119	1 1 2/1020000	0. 40
45	PLACE3000207	, , _,,,,	0. 32
15	PLACE3000230		0. 51
	PLACE3000271	F-PLACE3000271,	0. 41
	PLACE3000373	F-PLACE3000373	0. 37
	PLACE3000399	F-PLACE3000399	0. 42
20	PLACE3000401	F-PLACE3000401	0. 37
	PLACE3000406	F-PLACE3000406	0.46
	PLACE4000247	F-PLACE4000247	0. 64
	PLACE4000320	F-PLACE4000320	0. 36
25	PLACE4000344	F-PLACE4000344	0.31
	PLACE4000367	F-PLACE4000367	0.32
	PLACE4000401	F-PLACE4000401	0.54
	PLACE4000548	F-PLACE4000548	0. 26
30	THYR01000111	F-THYR01000111	0.53
	THYR01000129	F-THYR01000129	0. 45
	THYR01000187	F-THYR01000187	0.46
	THYR01000279	F-THYR01000279	0.09
35	THYR01000484	F-THYR01000484	0.36
	THYR01000596	F-THYR01000596	0.47
	THYR01000625	F-THYR01000625	0.59
•	THYR01000793	F-THYR01000793	0.17
40	THYR01000815	F-THYR01000815	0. 58
	THYR01000865	F-THYR01000865	0. 55
	THYR01001003	F-THYR01001003	0. 34
	THYR01001031	F-THYR01001031	0.43
45	THYR01001133	F-THYR01001133	0. 52
	THYR01001321	F-THYR01001321	0. 37
	THYR01001401		0. 36
	THYR01001426	F-THYR01001426	0. 32
50	THYR01001434		0.39
	THYR01001559		0. 45
	THYR01001570		0. 35
	THYR01001706		0.45
55	THYR01001746		
	THYR01001772	F-THYR01001772	0.37

Y79AA100033 F-Y79AA100033 0.53 Y79AA1000346 F-Y79AA1000346 0.53 Y79AA1000410 F-Y79AA1000410 0.13
Y79AA1000410 F-Y79AA1000410 0.13
1/3//1000410 1 1/3//1000410 0115
V7014400005 F V7014400005 0 22
Y79AA1000805 F-Y79AA1000805 0.32
Y79AA1001692 F-Y79AA1001692 0.45
Y79AA1002103 F-Y79AA1002103 0.15
Y79AA1002220 F-Y79AA1002220 0.49

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EXAMPLE 13

Full-length sequence analysis and homology search

Full-length sequence was determined for each selected cDNA clones. The nucleotide sequence determination was performed mainly by the dye-terminator method using custom synthesized DNA primers according to the primer walking procedure (custom synthesized DNA primers were used for sequencing; sequencing reaction was performed with DNA sequencing reagent supplied by PE Biosystems according to the supplier's manual; and the samples were analyzed in an automatic sequencer made by the same supplier). Sequence determination of some clones was carried out in the same manner but using a Licor DNA sequencer. Overlapping partial nucleotide sequences, which were obtained by the above-described method, were assembled together to determine a full-length nucleotide sequence. Amino acid sequences were then deduced from the determined full-length nucleotide sequences. However, amino acid sequence is not shown for a clone of which coding region was hard to be deduced or of which amino acid sequence has less than 100 amino acid residues. SEQ ID NOs corresponding to the respective clones are indicated in Table 350 and Table 351.

GenBank, Swiss-Prot and UniGene were searched for the determined nucleotide sequences by BLAST analysis. Matching data of cDNA clone which exhibits higher homology and of which functions are easily predicted based on the nucleotide sequences and the deduced amino acid sequences are selected from the BLAST analysis matching data with P value of 10⁻⁴à or less. The matching data selected are listed herein. The results of homology search 6, 12, 13, and 14 are indicated in the last part of this specification. However, there are some clones that did not match the criteria for judgment and such matching data of BLAST analysis are not shown herein.

EXAMPLE 14

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Novel full-length cDNA clone obtained from a cDNA library prepared by oligo-capping method

A cDNA clone, NT2RP4002298, was obtained from a cDNA library, NT2RP4 (see Example 1), prepared by oligo-capping method. Analysis of the entire nucleotide sequence of the clone has revealed that the clone encodes a novel protein consisting of 775 amino acids. The ATGpr1

score at the initiation codon of the amino acid sequence is 0.16, and therefore, the fullness ratio is low. However, the sequence can be full-length.

The full-length nucleotide sequence of NT2RP4002298 is shown in SEQ ID NO: 12370, and the deduced amino acid sequence encoded by the clone NT2RP4002298 is shown in SEQ ID NO: 12371.

EXAMPLE 15

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Gene expression analysis with hybridization using high density DNA filter

Nylon membrane for DNA spotting was prepared according to the following procedure. E. coli was cultured in each well of a 96-well plate (in a LB medium at 37°C for 16 hours). A small aliquot of each culture was suspended in 10 µl of sterile water in a well of a 96-well plate. The plate was heated at 100°C for 10 minutes. Then the boiled samples were analyzed by PCR reaction. PCR was performed in a 20 μl solution by using TaKaRa PCR Amplification Kit (Takara) according to the supplier's protocol. Primers used for the amplification of an insert cDNA in a plasmid were a pair of sequencing primers, ME761FW (5' tacggaagtgttacttctgc 3' / SEQ ID NO: 13290) and ME1250RV (5' tgtgggaggttttttctcta 3' / SEQ ID NO: 13291), or a pair of primers, M13M4 (5' gttttcccagtcacgac 3' / SEQ ID NO: 13292) and M13RV (5' caggaaacagctatgac 3' / SEQ ID NO: 13293). PCR reaction was performed in a thermal cycler, GeneAmp System 9600 (PE Biosystems). The cycling profile consisted of pre-heat at 95°C for 5 mimutes; 10 cycles of denaturation at 95°C for 10 seconds, and annealing/extension at 68°C for 1 minute; 20 cycles of denaturation at 98°C for 20 seconds and annealing/extension at 60°C for 3 minutes; and final extension at 72°C for 10 minutes. After the PCR reaction, the 20 µl reaction solution was loaded onto a 1 % agarose gel and fractionated by electrophoresis. DNA on the gel was stained with ethidium bromide to confirm the amplification of cDNA. When cDNAs were barely amplified by PCR, plasmids containing the corresponding insert cDNAs were prepared by the alkali-extraction method (J. Sambrook, E.F., Fritsh, & T. Maniatis, "Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989).

Preparation of DNA array was carried out by the following procedure. An Aliquot of a DNA solution was added in each well of a 384-well plate. DNA was spotted onto a nylon membrane (Boehringer) by using a 384-pin tool of Biomek 2000 Laboratory Automation Sysytem (Beckman-Coulter). Specifically, the 384-well plate containing the DNA was placed under the 384-pin tool. The independent 384 needles were simultaneously dipped into the DNA solution for DNA deposition. The needles were gently pressed onto a nylon membrane and the DNA deposited at the tips of needles was spotted onto the membrane. Denaturation of the spotted DNA and immobilization of the DNA on the nylon membrane were carried out according to usual manners (J. Sambrook, E.F., Fritsh, & T. Maniatis, "Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989).

Hybridization probe used was radioisotope-labeled 1st strand cDNA. The 1st strand cDNA synthesis was performed by using Thermoscript^(TM) à RT-PCR System (GIBCO). Specifically, the 1st strand cDNA was synthesized by using 1.5 μ g mRNAs from various human tissues (Clontech), 1 μ l aliquots of 50 μ M Oligo(dT)20 and 50 μ Ci [α ³³àP]dATP according to an attached protocol. Probe purification was carried out by using ProbeQuant^(TM) à G-50 micro column (Amersham-Pharmacia Biotech) according to an attached protocol. In the next step, 2 units of E.

coli RNasH were added to the reaction mixture. The mixture was incubated at room temperature for 10 minutes and then 100 μg of human COT-1 DNA (GIBCO) was added thereto. The mixture was incubated at 97°C for 10 minutes and then was allowed to stand on ice to give hybridization probe.

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Hybridization of the radioisotope-labeled probe to the DNA array was performed in a usual manner (J. Sambrook, E.F., Fritsh, & T. Maniatis, Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989). The membrane was washed as follows: the nylon membrane was incubated in Washing solution 1 (2 x SSC, 1% SDS) at room temperature (about 26°C) for 20 minutes and this washing was repeated 3 times; then the membrane was washed 3 times by incubating it in Washing solution 2 (0.1 x SSC, 1% SDS) at 65°C for 20 minutes. Autoradiography was performed by using an image plate for BAS2000 (Fuji Photo Film Co., Ltd.). Specifically, the nylon membrane with probe hybridized thereon was wrapped with a piece of Saran Wrap and brought into tight contact with the image plate on the light-sensitive surface. The membrane with the image plate was placed in an imaging cassette for radioisotope and allowed to stand in dark place for 4 hours. The radioactivity recorded on the image plate was analyzed by using BAS2000 (Fuji Photo Film Co., Ltd.). The activity was subjected to electronic conversion and recorded as an image file of autoradiogram. The signal intensity of each DNA spot was analyzed by using Visage High Density Grid Analysis Systems (Genomic Solutions Inc.). The signal intensity was converted into numerical data. The data were taken by duplicated measurements. The reproducibility was assessed by comparing the signal intensities of the corresponding spots on the duplicated DNA filters that were hybridized to a single DNA probe (Figure 2). The ratio between the corresponding spots falls within a range of 2 or less in 95% of entire spots and the correlation coefficient is r=0.97. Thus the reproducibility is assumed to be satisfactory.

The detection sensitivity in gene expression analysis was estimated by examining increases in the signal intensity of probe concentration-dependent spot in hybridization using a probe complementary to the DNA spotted on the nylon membrane. DNA used was PLACE1008092 (same as DNA deposited in GenBank under an Accession No. AF107253). The DNA array with DNA of PLACE1008092 was prepared according to the above-mentioned method. The probe used was prepared as follows: mRNA was synthesized in vitro from the clone, PLACE1008092. By using this mRNA as a template, radioisotope-labeled 1st strand cDNA was synthesized in the same manner as described above, and the cDNA was used as the probe. The cDNA PLACE1008092 was inserted into pBluescript SK(-), of which T7 promoter was ligated to the 5' end of the cDNA, to give a plasmid for in vitro synthesis of the mRNA from PLACE1008092. Specifically, the PLACE1008092 insert was cut out from pME18SFL3 carrying the cDNA at a DrallI site thereof by Xhol digestion. The resulting PLACE1008092 fragment was ligated to Xholpredigested pBluescript SK(-) by using DNA ligation kit ver.2 (Takara). The in-vitro mRNA synthesis from PLACE1008092 inserted in pBluescript SK(-) was carried out by using Ampliscribe (TM) à T7 high yield transcription kit (Epicentre technologies). Hybridization and the analysis of signal intensity of each DNA spot were conducted by the same methods as described above. When the probe concentration is 1×10^{7} à µg/ml or less, there was no increase of signal intensity proportional to the probe concentration. Therefore it was assumed to be difficult to compare the signals with one another in the concentration range. Thus the spots with the intensity of 40 or less were indiscriminately taken as low-level signals (Figure 3). Within a concentration of the probe ranging from $1 \times 10^7 \text{à} \, \mu\text{g/m}$ to 0.1 $\mu\text{g/m}$, the signal was

found to increase in a probe concentration-dependent manner. The detection limit was then assumed to be 1:100,000 in a ratio of mRNA expression level in a sample.

Tables 12-168 (also containing clones with no description in Examples) show the expression of each cDNA in human normal tissues (heart, lung, pituitary gland, thymus, brain, kidney, liver and spleen). The expression levels are indicated by numerical values of 0-10,000. Genes that were expressed in at least a single tissue are indicated below by the corresponding clone names:

10	HEMBA1000012,	HEMBA1000020,	HEMBA1000030,	HEMBA1000042,	HEMBA1000046,
	HEMBA1000076,	HEMBA1000111,	HEMBA1000129,	HEMBA1000150,	HEMBA1000156,
	HEMBA1000158,	HEMBA1000168,	HEMBA1000185,	HEMBA1000216,	HEMBA1000227,
15	HEMBA1000231,	HEMBA1000243,	HEMBA1000244,	HEMBA1000280,	HEMBA1000282,
15	HEMBA1000288,	HEMBA1000303,	HEMBA1000304,	HEMBA1000327,	HEMBA1000338,
	HEMBA1000351,	HEMBA1000355,	HEMBA1000356,	HEMBA1000357,	HEMBA1000366,
	HEMBA1000369,	HEMBA1000376,	HEMBA1000387,	HEMBA1000392,	HEMBA1000396,
20	HEMBA1000411,	HEMBA1000422,	HEMBA1000428,	HEMBA1000456,	HEMBA1000459,
	HEMBA1000469,	HEMBA1000488,	HEMBA1000491,	HEMBA1000501,	HEMBA1000505,
	HEMBA1000508,	HEMBA1000519,	HEMBA1000523,	HEMBA1000534,	HEMBA1000540,
	HEMBA1000542,	HEMBA1000545,	HEMBA1000557,	HEMBA1000561,	HEMBA1000568,
25	HEMBA1000569.	HEMBA1000575,	HEMBA1000588,	HEMBA1000591.	HEMBA1000604.

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			EP 1 0/4 61/	AZ	
	-HEMBA1000622,	HEMBA1000636,	HEMBA1000655,	HEMBA1000657,	HEMBA1000673,
	HEMBA1000682,	HEMBA1000702,	HEMBA1000705,	HEMBA1000726,	HEMBA1000749,
	HEMBA1000752,	HEMBA1000769,	HEMBA1000774,	HEMBA1000791,	HEMBA1000827,
5	HEMBA1000843,	HEMBA1000852,	HEMBA1000870,	HEMBA1000872,	HEMBA1000876,
	HEMBA1000908,	HEMBA1000910,	HEMBA1000918,	HEMBA1000919,	HEMBA1000934,
	HEMBA1000942,	HEMBA1000960,	HEMBA1000971.	HEMBA1000972,	HEMBA1000975,
	HEMBA1000986,	HEMBA1000991,	HEMBA1001007,	HEMBA1001008,	HEMBA1001017,
10	HEMBA1001020,		-	MBA1001060, HEMBA1	•
	HEMBA1001080,	HEMBA1001085,	HEMBA1001088,	1177777 4 4 4 4 4 4 4 4 4 4 4 4 4	HEMBA1001121,
	HEMBA1001123,	HEMBA1001133,	HEMBA1001137,		HEMBA1001172,
	HEMBA1001120,	HEMBA1001208,	HEMBA1001213,	1100011001000	HEMBA1001235,
15	HEMBA1001265,	HEMBA1001281,	HEMBA1001286,	HEMBA1001289,	HEMBA1001294.
,-	HEMBA1001299,	HEMBA1001302,	HEMBA1001303,	HEMBA1001310,	НЕМВА1001326,
	HEMBA1001330,	HEMBA1001351,	HEMBA1001375,	HEMBA1001377,	HEMBA1001387,
	HEMBA1001388,	HEMBA1001398,	HEMBA1001405,		HEMBA1001415,
20		HEMBA1001333,	HEMBA1001435,		HEMBA1001450,
20	HEMBA1001432,	HEMBA1001463,	HEMBA1001476,		HEMBA1001510-
	HEMBA1001454, HEMBA1001515,	HEMBA1001517,	HEMBA1001526,		HEMBA1001557,
	HEMBA1001516,	HEMBA1001569,	HEMBA1001570,		HEMBA1001581,
05	HEMBA1001589,	HEMBA1001595,	HEMBA1001608,		HEMBA1001636;
25	HEMBA1001640.	HEMBA1001647,	HEMBA1001651,	************	HEMBA1001661,
	HEMBA1001678,	HEMBA1001709,	HEMBA1001712,		HEMBA1001718,
	HEMBA1001731,	HEMBA1001734,	HEMBA1001745.		HEMBA1001781,
	HEMBA1001784,	HEMBA1001791,	HEMBA1001804,		HEMBA1001809,
30	HEMBA1001815,	HEMBA1001822,	HEMBA1001824,		HEMBA1001847.
	HEMBA1001864,	HEMBA1001866,	HEMBA1001869,	III) III . 4 6 6 4 6 6 6	HEMBA1001896,
	HEMBA1001910,	HEMBA1001912,	HEMBA1001913,		HEMBA1001940,
	HEMBA1001964,	•		MBA1001991, HEMBA1	•
35	HEMBA1002008,	HEMBA1002018,	HEMBA1002035,		HEMBA1002092,
	HEMBA1002100,	HEMBA1002102,	HEMBA1002113		HEMBA1002125,
	HEMBA1002144,	HEMBA1002150,	HEMBA1002151,		HEMBA1002161,
	HEMBA1002162,	HEMBA1002166,	HEMBA1002177		HEMBA1002189,
40	HEMBA1002191,	HEMBA1002226,	HEMBA1002229		HEMBA1002241,
	HEMBA1002265,	HEMBA1002267,	HEMBA1002270		HEMBA1002337.
	HEMBA1002341,	HEMBA1002363,	HEMBA1002381		HEMBA1002417,
	HEMBA1002419,	HEMBA1002439,	HEMBA1002458		HEMBA1002469,
45	HEMBA1002477,	HEMBA1002486,	HEMBA1002495		HEMBA1002503,
	HEMBA1002508,	HEMBA1002513,	HEMBA1002515	·	HEMBA1002542,
	HEMBA1002552,	HEMBA1002558,	HEMBA1002561	·	HEMBA1002590,
	HEMBA1002592,	HEMBA1002609,	HEMBA1002624		HEMBA1002645,
50	HEMBA1002659,	HEMBA1002661,	HEMBA1002678		HEMBA1002696,
	HEMBA1002703,	HEMBA1002712,	HEMBA1002728		HEMBA1002746,
	HEMBA1002748,		HEMBA1002768		HEMBA1002779.
	HEMBA1002780,	HEMBA1002794,	HEMBA1002810		HEMBA1002818,
55	HEMBA1002826,	HEMBA1002833,	HEMBA1002863	•	HEMBA1002896,
	HEMBA1002921,	HEMBA1002924,	HEMBA1002934		HEMBA1002944,

HEMBA10032968,				EP 1 0/4 61/ A	2	
HEMBA1003046, HEMBA1003067, HEMBA1003071, HEMBA1003078, HEMBA1003079, HEMBA1003083, HEMBA1003124, HEMBA1003124, HEMBA1003124, HEMBA1003129, HEMBA1003127, HEMBA1003199, HEMBA1003204, HEMBA1003212, HEMBA1003226, HEMBA1003257, HEMBA1003227, HEMBA1003226, HEMBA1003257, HEMBA1003228, HEMBA1003228, HEMBA1003288, HEMBA1003204, HEMBA1003330, HEMBA1003348, HEMBA1003370, HEMBA1003373, HEMBA1003376, HEMBA1003380, HEMBA1003384, HEMBA1003403, HEMBA1003373, HEMBA1003418, HEMBA1003380, HEMBA1003561, HEMBA1003565, HEMBA1003561, HEMBA1003561, HEMBA1003661, HEMBA1003667, HEMBA1003677, HEMBA1003667, HEMBA1003675, HEMBA1003667, HEMBA1003675, HEMBA1003666, HEMBA1003666, HEBBA1003666, HEBBA1003666, HEMBA1003666,		HEMBA1002968,	HEMBA1002970,	HEMBA1002971,	HEMBA1002973,	HEMBA1002997,
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HEMBA1003136, HEMBA1003142, HEMBA1003148, HEMBA1003216, HEMBA1003175, HEMBA1003179, HEMBA1003229, HEMBA1003220, HEMBA1003220, HEMBA1003220, HEMBA1003221, HEMBA1003257, HEMBA1003273, HEMBA1003278, HEMBA1003278, HEMBA1003281, HEMBA1003286, HEMBA1003286, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003330, HEMBA1003348, HEMBA1003348, HEMBA1003370, HEMBA1003373, HEMBA1003376, HEMBA1003358, HEMBA1003408, HEMBA1003376, HEMBA1003538, HEMBA1003408, HEMBA1003481, HEMBA1003528, HEMBA1003558, HEMBA1003558, HEMBA1003558, HEMBA1003559, HEMBA1003556, HEMBA1003559, HEMBA1003557, HEMBA1003559, HEMBA1003557, HEMBA1003559, HEMBA1003557, HEMBA1003559, HEMBA1003557, HEMBA1003559, HEMBA1003572, HEMBA1003559,		HEMBA1003046,	HEMBA1003067,	HEMBA1003071,	HEMBA1003078,	HEMBA1003079,
HEMBA1003136, HEMBA1003142, HEMBA1003148, HEMBA1003216, HEMBA1003175, HEMBA1003179, HEMBA1003229, HEMBA1003220, HEMBA1003220, HEMBA1003220, HEMBA1003221, HEMBA1003257, HEMBA1003273, HEMBA1003278, HEMBA1003278, HEMBA1003281, HEMBA1003286, HEMBA1003286, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003328, HEMBA1003330, HEMBA1003348, HEMBA1003348, HEMBA1003370, HEMBA1003373, HEMBA1003376, HEMBA1003358, HEMBA1003408, HEMBA1003376, HEMBA1003538, HEMBA1003408, HEMBA1003481, HEMBA1003528, HEMBA1003558, HEMBA1003558, HEMBA1003558, HEMBA1003559, HEMBA1003556, HEMBA1003559, HEMBA1003557, HEMBA1003559, HEMBA1003557, HEMBA1003559, HEMBA1003557, HEMBA1003559, HEMBA1003557, HEMBA1003559, HEMBA1003572, HEMBA1003559,	5	HEMBA1003083,	HEMBA1003086,	HEMBA1003098,	HEMBA1003129,	HEMBA1003133.
HEMBA1003179, HEMBA1003199, HEMBA1003202, HEMBA1003204, HEMBA1003212, HEMBA1003227, HEMBA1003227, HEMBA1003227, HEMBA1003227, HEMBA1003227, HEMBA1003281, HEMBA1003328, HEMBA1003329, HEMBA1003328, HEMBA1003330, HEMBA1003348, HEMBA1003328, HEMBA1003330, HEMBA1003348, HEMBA1003348, HEMBA1003348, HEMBA1003403, HEMBA1003408, HEMBA1003418, HEMBA1003418, HEMBA1003511, HEMBA1003555, HEMBA1003556, HEMBA1003550, HEMBA1003551, HEMBA1003559, HEMBA1003557, HEMBA1003559, HEMBA1003557, HEMBA1003551,				HEMBA1003148.	HEMBA1003166.	
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Genes that were expressed in all the tissues tested are indicated below by the corresponding clone names:

	HEMBA1000020,	HEMBA1000150,	HEMBA1000282,	HEMBA1000376,	HEMBA1000387,
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55	PLACE1001761,	PLACE1001912,	PLACE1002004,	PLACE1002066,	PLACE1002532,
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	_		EP 1 074 617 AZ	2	
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35	Y79AA1002482.				

Genes that were expressed at low levels in any of the tissues tested are indicated below by the corresponding clone names:

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	HEMBA1000302,	HEMBA1000307,	HEMBA1000333,	HEMBA1000390,	HEMBA1000418,
5	HEMBA1000434,	HEMBA1000442,	HEMBA1000460,	HEMBA1000464,	HEMBA1000490,
	HEMBA1000504,	HEMBA1000518,	HEMBA1000520,	HEMBA1000531,	HEMBA1000555,
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	HEMBA1000662,	HEMBA1000686,	HEMBA1000719,	HEMBA1000722,	HEMBA1000727,
10	HEMBA1000747,	HEMBA1000773,	HEMBA1000817,	HEMBA1000822,	HEMBA1000851,
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45	Y79AA1002487.				

Genes exhibiting characteristic features in the expression thereof were selected by statistical analysis of these data. Two examples are shown below to describe the selection of genes of which expression is varied greatly among tissues. The β ' actin gene is used frequently as a control in gene expression analysis. Genes of which expression is varied greatly among tissues as compared that of the β -actin gene were determined as follows. Specifically, sum of squared deviation was calculated in the signal intensity of β -actin observed in each tissue, which was divided by 7 degrees of freedom to determine a variance $S_a^{\ 2}$ à. Next, sum of squared deviation was calculated in the signal intensity of a gene to be compared observed in

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each tissue, which was divided by 7 degrees of freedom to determine a variance $S_b^2 \dot{a}$. By taking variance ratio F as $F = S_b^2 \dot{a}/S_a^2 \dot{a}$, genes with a significance level of 5% or more were extracted in the F distribution. Genes extracted are indicated below by the corresponding clone names:

HEMBA1002113, HEMBA1005296, HEMBA1007121, HEMBB1000637, HEMBB1000915, MAMMA1000597, MAMMA1000605, MAMMA1000962, MAMMA1001139, MAMMA1001198, MAMMA1002858, NT2RM2001896, NT2RP2002710, NT2RP2004339, NT2RP2004538, NT2RP3000348, NT2RP3003121, PLACE3000009, PLACE3000254, THYRO1000569, Y79AA1000131.

Gene of OVARC1000037 {heterogeneous nuclear ribonucleoprotein (hnRNP)} which expression is varied little. Genes of which expression is varied greatly among tissues as compared that of the OVARC1000037 gene were determined as follows. Specifically, sum of squared deviation was calculated in the signal intensity of β -actin observed in each tissue, which was divided by 7 degrees of freedom to determine a variance S_a^2 à. Next, sum of squared deviation was calculated in the signal intensity of a gene to be compared observed in each tissue, which was divided by 7 degrees of freedom to determine a variance S_b^2 à. By taking variance ratio F as $F = S_b^2$ à/ S_a^2 à, genes with a significance level of 5% or more were extracted in the F distribution. Genes extracted are indicated below,by the corresponding clone names:

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NT2RM2001716,	NT2RM2001840,
HEMBA1000726,	HEMBA1001286,

			LI I UI T UII AL		
	HEMBA1000304,	NT2RM2001716,	NT2RM2001840,	HEMBA1001051,	HEMBA1001109,
	OVARC1001731,	HEMBA1000726,	HEMBA1001286,	HEMBA1000387,	HEMBA1000519,
	NT2RM2001896,	-HEMBA1000042,	HEMBA1001085,	HEMBA1001330,	0VARC1000576_
5	HEMBA1000575,	NT2RM2000599,	NT2RM2000714,	HEMBA1000469,	NT2RM4000366,
	HEMBA1001377,	HEMBA1000769,	HEMBA1000338,	NT2RM2000795,	HEMBA1001299,
	HEMBA1000508,	HEMBA1000150,	HEMBA1000774,	HEMBA1001226,	HEMBA1000960;
	NT2RM4000795,	HEMBA1002162,	NT2RM4001876,	NT2RM4002482,	HEMBA1001678,
10	HEMBA1002113,	NT2RM4002383,	HEMBA1002229,	HEMBA1002818,	HEMBA1001454,
	NT2RM4000764,	HEMBA1001510,	HEMBA1001714,	HEMBA1002150,	NT2RM4002044,
	HEMBA1002728,	NT2RM4002189,	HEMBA1001991,	HEMBA1002166,	NT2RM4002499,
	NT2RM4001140,	NT2RM4002504,	HEMBA1002590,	HEMBA1001435,	PLACE1000706,
15	HEMBA1002160,	HEMBA1001824,	HEMBA1001463,	HEMBA1001533,	HEMBA1001570,
	PLACE1001036,	HEMBA1001651,	HEMBA1002381,	HEMBA1002934,	HEMBA1003370,
	HEMBA1003021,	HEMBA1003166,	NT2RP1000738,	NT2RP2000040,	HEMBA1004164,
	HEMBA1003836,	HEMBA1004267,	NT2RP2000845,	HEMBA1003041,	HEMBA1003571,
20	HEMBA1003758,	NT2RP2000108,	HEMBA1003838,	NT2RP1000357,	HEMBA1003376,
	PLACE1003528,	HEMBA1003528,	NT2RP1001475,	HEMBA1004049,	HEMBA1003212,
	HEMBA1003667,	PLACE1004149,	HEMBA1003926,	HEMBA1004306,	HEMBA1004024,
	NT2RP1000363,	HEMBA1003033,	HEMBA1004335,	HEMBA1003348,	HEMBA1003034,
25	NT2RP2001081,	немва1004056, не	MBA1003314, HEMB.	A1003827, HEMBA1	003893,
	NT2RP2001036,	NT2RP2001168,	NT2RP2001328,	HEMBA1005035,	NT2RP2001569,
	NT2RP2002439,	HEMBA1005511,	HEMBA1005999,	NT2RP2002862,	NT2RP2002979,
	NT2RP2001394,	HEMBA1004753,	NT2RP2002621,	HEMBA1005853,	HEMBA1005443,
30	NT2RP2002980,	NT2RP2001347,	HEMBA1005241,	NT2RP2002750,	NT2RP2003533,
	HEMBA1005634,	NT2RP2003034,	HEMBA1006138,	NT2RP2003117,	NT2RP2001366,
	HEMBA1005079,	NT2RP2003293,	NT2RP2002710,	HEMBA1005911,	NT2RP2002752,
35	HEMBA1006036,	NT2RP2002987,	HEMBA1006100,	HEMBA1004460,	HEMBA1004538,
33	NT2RP2001943,	NT2RP2002033,	HEMBA1005296,	HEMBA1005829,	HEMBA1005520,
	HEMBA1005123,	HEMBA1005552,	HEMBA1004930,	NT2RP2001312,	HEMBA1005304,
	HEMBA1005834,	HEMBA1005990,	HEMBA1005526,	NT2RP2003073,	HEMBA1005331,
40	HEMBA1006744,	HEMBA1006780,	NT2RP2004339,	HEMBB1000173,	HEMBA1007113,
,,	NT2RP2005908,	HEMBB1000376,	HEMBB1000024,	HEMBB1000510,	NT2RP2004580,

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	NT2RP2003912,	HEMBA1006381,	NT2RP2004270,	HEMBA1006993,	NT2RP2005325,
	HEMBB1000218,	HEMBA1007121,	HEMBB1000434,	NT2RP2006166,	PLACE1010310,
	HEMBB1000589,	NT2RP2003986,	HEMBA1006419,	HEMBB1000141,	HEMBB1000030,
5	HEMBB1000099,	HEMBA1006624,	NT2RP2003988,	NT2RP2005753,	HEMBA1007147,
	HEMBB1000354,	NT2RP2006023,	HEMBB1000441,	HEMBB1000523,	HEMBB1000059,
	HEMBA1006631,	NT2RP2004538,	HEMBA1006695,	HEMBB1000490,	NT2RP2006184,
	HEMBB1000573,	NT2RP2004675,	HEMBA1007078,	NT2RP2005457,	NT2RP2005491,
10	HEMBB1000491,	HEMBB1000575, PLA	•	2005773, NT2RP20	•
	HEMBB1000337,	HEMBB1000008,	NT2RP2005701,	NT2RP2003825,	HEMBB1000215,
	HEMBB1000258,	NT2RP2005496,	HEMBB1000374,	NT2RP2005581,	HEMBB1000018,
	HEMBB1000493,	HEMBB1000554,	HEMBB1000671,	NT2RP3000348,	HEMBB1001267,
15	HEMBB1002092,	HEMBB1001749.	PLACE3000304,	HEMBB1001834,	HEMBB1001142.
,,,	HEMBB1001177,	HEMBB1000840,	HEMBB1001436,	HEMBB1001921,	HEMBB1002005,
	NT2RP3001398,	HEMBB1002069,	HEMBB1002094,	HEMBB1001836,	HEMBB1001114,
	NT2RP3000628,	HEMBB1001367,	HEMBB1001588,	HEMBB1002249,	HEMBB1000684,
20	NT2RP2006571,	HEMBB1001802,	HEMBB1000915,	HEMBB1000996,	NT2RP3001274,
20	HEMBB1001564,	NT2RP3000968,	PLACE3000156,	HEMBB1002045,	NT2RP3000341
	HEMBB1000917,	PLACE1011090,	NT2RP3000742.	PLACE2000435,	HEMBB1001944.
	PLACE3000254,	HEMBB1002218,	NT2RP2006436,	NT2RP3000320,	HEMBB1001126,
	HEMBB1000749,	NT2RP3000644,	HEMBB1001527,	NT2RP3000850,	HEMBB1001967;
25	PLACE3000401,	HEMBB1000637,	HEMBB1001234,	NT2RP3000080,	PLACE2000411,
	PLACE3000009,	PLACE3000070,	HEMBB1002134,	PLACE3000475,	PLACE4000192,
	HEMBB1002520,	HEMBB1002545,	MAMMA1000155,	MAMMA1000307,	THYR01000787,
	MAMMA1000348,	MAMMA1000372,	NT2RP3003121,	MAMMA1000851,	MAMMA1000501,
30	THYR01000569,	MAMMA1000576,	MAMMA1000605,	HEMBB1002442,	HEMBB1002596,
	MAMMA1000198,	MAMMA1000714,	MAMMA1000356,	MAMMA1000760,	MAMMA1000416,
	MAMMA1000931,	MAMMA1000962,	HEMBB1002387,	NT2RP3001712,	HEMBB1002556,
	HEMBB1002617,	MAMMA1000171, PLA	CE4000612, MAMMA	A1000761, MAMMA10	000421,
35	MAMMA1000444,	HEMBB1002358,	HEMBB1002453,	HEMBB1002664,	HEMBB1002699,
	MAMMA1000720,	MAMMA1000360,	MAMMA1000744,	MAMMA1000802,	MAMMA1000839,
	MAMMA1000478,	MAMMA1000877,	MAMMA1000585,	MAMMA1000941,	NT2RP3002909,
	MAMMA1000968,	HEMBB1002457,	HEMBB1002495,	MAMMA1000057,	MAMMA1000696,
40	иамма 1000942,	MAMMA1000302,	MAMMA1000943,	MAMMA1000998,	NT2RP3001752,
	NT2RP3003032,	MAMMA1000429,	MAMMA1000565,	MAMMA1000594,	MAMMA1000625,
	NT2RP3002146,	MAMMA1000084,	MAMMA1000257,	NT2RP3003157,	MAMMA1000431,
	MAMMA1000597,	MAMMA1001078,	MAMMA1001126,	THYR01001537,	NT2RP3004093,
45	MAMMA1001202,	NT2RP3003842,	Y79AA1000342,	MAMMA1002311,	MAMMA1001969,
	Y79AA1000560,	MAMMA1002359,	MAMMA1002056,	MAMMA1002413,	MAMMA1002454,
	MAMMA1002612,	MAMMA1001411,	MAMMA1001035,	MAMMA1001080,	MAMMA1001133,
	NT2RP3004095,	MAMMA1001161,	MAMMA1001203,	MAMMA1001663,	NT2RP3003819,
50	MAMMA1001745,	MAMMA1001790,	NT2RP3004470,	NT2RP3004503,	MAMMA1002293,
	NT2RP4000321,	MAMMA1001880,	Y79AA1000410,	MAMMA1001970,	MAMMA1002058,
	NT2RP4000515,	MAMMA1002215,	MAMMA1002617,	NT2RP3003576,	MAMMA1001139,
	MAMMA1001576,	Y79AA1000131,	MAMMA1001330,	MAMMA1001769,	MAMMA1001383,
55	MAMMA1002032,	MAMMA1002125,	MAMMA1002174,	NT2RP4000614,	NT2RP4000243,
	NT2RP3004110,	MAMMA1001271,	MAMMA1001296,	MAMMA1001992,	MAMMA1002033,

	MAMMA1002428,	MAMMA1002590,	MAMMA1001186,	MAMMA1002267,	MAMMA1002322,
	MAMMA1001956,	MAMMA1002155, NT2	RP4000210, MAMMA	1002622, NT2RP3	004125,
	MAMMA1001220,	MAMMA1001683,	NT2RP3004348,	Y79AA1000214,	Y79AA1000833,
5	NT2RP4000212,	MAMMA1002230,	MAMMA1001452,	MAMMA1001620,	MAMMA1001256,
	MAMMA1001760,	NT2RP3004349,	MAMMA1001783,	MAMMA1001907,	MAMMA1002009,
	MAMMA1002545,	NT2RP4000214,	NT2RP4000728,	MAMMA1001465,	MAMMA1001154,
	MAMMA1001198,	MAMMA1001343,	MAMMA1002310,	NT2RP4000035,	NT2RP4000833,
10	MAMMA1003150,	MAMMA1002886,	NT2RP4001938,	NT2RM2000260,	MAMMA1002629,
	MAMMA1002973,	MAMMA1002721,	MAMMA1002909,	NT2RP4001100,	NT2RM1000857,
	NT2RP4000878,	MAMMA1002844,	NT2RM1000039,	NT2RP4001174,	MAMMA1002665,
	MAMMA1003047,	NT2RM1000086,	NT2RM1000260,	NT2RM1000355,	MAMMA1002701,
15	NT2RP4000918,	MAMMA1002830,	MAMMA1002970,	NT2RP4001677,	NT2RM2000422,
	MAMMA1003004,	MAMMA1002673,	MAMMA1003031,	MAMMA1002764,	MAMMA1002858,
	NT2RP4001679,	NT2RP4002888,	MAMMA1002711,	NT2RP4001276,	NT2RM1000018,
20	NT2RP4001568,	NT2RM1000883.			

Thus, characteristic features in the expression of a gene are illustrated by comparing and _ statistically analyzing the expression of many genes.

Analysis of disease-associated genes

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Non-enzymic protein glycation reaction is believed to be a cause of a variety of chronic diabetic complications. Accordingly, genes of which expression is elevated or decreased in a glycated protein-specific manner in the endothelial cells are associated with diabetic complications caused by glycated proteins. Vascular endothelial cells are affected with glycated proteins present in blood. Reaction products of non-enzymic protein glycation include amadori compound (glycated protein) as a mildly glycated protein and advanced glycation endproduct as a heavily glycated protein. Hence, a survey was carried out for genes of which expression levels are varied depending on the presence of these glycated proteins in endothelial cells. The mRNAs were extracted from endothelial cells that were cultured in the presence or absence of glycated protein. The mRNAs were converted into radiolabeled first strand cDNAs for preparing probes. The probes were hybridized to the above-mentioned DNA array. Signal of each DNA spot was detected by BAS2000 and analyzed by ArrayGauge (Fuji Photo Film Co., Ltd.).

Advanced glycation endproduct of bovine serum albumin was prepared as follows: bovine serum albumin (BSA; Sigma) was incubated in a phosphate buffer solution containing 50 mM glucose at 37°C for 8 weeks; and the resulting brownish BSA was dialyzed against a phosphate buffer solution.

Human normal pulmonary arterial endothelial cells (Cell Applications) were cultured in an Endothelial Cell Growth Medium (Cell Applications). The culture dish (Farcon) with the cells were incubated in a CO₂ incubator (37°C, 5% CO₂, in a humid atmosphere). When the cells were grown to be confluent in the dish, 250 μg/ml of bovine serum albumin (sigma), glycated bovine serum albumin (Sigma) or advanced glycation endproduct of bovine serum albumin was added thereto and the cells were incubated for 33 hours. The mRNA was extracted from the cells by using a FastTack^(TM) à 2.0 kit (Invitrogen). The labeling of hybridization probe was carried out

by using the mRNA according to the same procedure as described above.

Table 169 shows the expression level of each cDNA in human pulmonary arterial endothelial cells cultured in a medium containing bovine serum albumin (sigma), glycated bovine serum albumin (Sigma) or advanced glycation endproduct of bovine serum albumin. Genes of which expression was detected in the endothelial cell are as follows:

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				14.	
	HEMBA1000005,	HEMBA1000012,	HEMBA1000020,	HEMBA1000042,	HEMBA1000046,
	HEMBA1000076,	HEMBA1000111,	HEMBA1000150,	HEMBA1000185,	HEMBA1000216.
	HEMBA1000231,	HEMBA1000280,	HEMBA1000282,	HEMBA1000288,	HEMBA1000304,
5	HEMBA1000327,	HEMBA1000338,	HEMBA1000356,	HEMBA1000357,	HEMBA1000366.
	HEMBA1000376,	HEMBA1000387,	HEMBA1000428,	HEMBA1000456,	HEMBA1000459.
	HEMBA1000460,	HEMBA1000469,	HEMBA1000501,	HEMBA1000504,	***************************************
	HEMBA1000508,	HEMBA1000519,			HEMBA1000505,
			HEMBA1000534,	HEMBA1000540,	HEMBA1000542,
10	HEMBA1000555,	HEMBA1000557,	HEMBA1000575,	HEMBA1000592,	HEMBA1000604,
	HEMBA1000622,	HEMBA1000636,	HEMBA1000655,	HEMBA1000673,	HEMBA1000682,
	HEMBA1000686,	HEMBA1000702,	HEMBA1000726,	HEMBA1000727,	HEMBA1000749,
	HEMBA1000769,	HEMBA1000774,	HEMBA1000791,	HEMBA1000827,	HEMBA1000852,
15	HEMBA1000872,	HEMBA1000876,	HEMBA1000918,	HEMBA1000942,	HEMBA1000960,
	HEMBA1000974,	HEMBA1001020,	HEMBA1001051,	HEMBA1001059,	HEMBA1001060,
	HEMBA1001080,	HEMBA1001085,	HEMBA1001088,	HEMBA1001109,	HEMBA1001122,
	HEMBA1001123,	HEMBA1001140,	HEMBA1001172,	HEMBA1001196,	HEMBA1001197,
20	HEMBA1001213,	HEMBA1001226,	HEMBA1001235,	HEMBA1001247,	HEMBA1001265,
	HEMBA1001281,	HEMBA1001286,	HEMBA1001299,	HEMBA1001302,	HEMBA1001326,
	HEMBA1001330,	HEMBA1001351,	HEMBA1001377,	HEMBA1001387,	HEMBA1001398,
	HEMBA1001432,	HEMBA1001433,	HEMBA1001435,	HEMBA1001446,	HEMBA1001450,
25	HEMBA1001454,	HEMBA1001463, HEM	BA1001497, HEMB	A1001510, HEMBA1	001517,
25	HEMBA1001526,	HEMBA1001533,	HEMBA1001557,	HEMBA1001566,	HEMBA1001569,
į	HEMBA1001570,	HEMBA1001579,	HEMBA1001581,	HEMBA1001595,	HEMBA1001608,
	HEMBA1001620,	HEMBA1001640,	HEMBA1001647,	HEMBA1001651,	HEMBA1001655,
	HEMBA1001658,	HEMBA1001678,	HEMBA1001714,	HEMBA1001718,	HEMBA1001723,
30	HEMBA1001746,	HEMBA1001784,	HEMBA1001791,	HEMBA1001800,	HEMBA1001804,
	HEMBA1001809,	HEMBA1001819,	HEMBA1001824,	HEMBA1001844,	HEMBA1001866,
	HEMBA1001888,	HEMBA1001912,	HEMBA1001913,	HEMBA1001918,	HEMBA1001940,
	HEMBA1001960,	HEMBA1001987,	HEMBA1001991,	HEMBA1002003,	HEMBA1002008,
35	HEMBA1002049,	HEMBA1002100,	HEMBA1002102,	HEMBA1002113,	HEMBA1002119,
	HEMBA1002125,	HEMBA1002151,	HEMBA1002153,	HEMBA1002160,	HEMBA1002161,
	HEMBA1002166,	HEMBA1002185,	HEMBA1002189,	HEMBA1002191,	HEMBA1002226,
	HEMBA1002229,	HEMBA1002237,	HEMBA1002241,	HEMBA1002267,	HEMBA1002270,
40	HEMBA1002337,	HEMBA1002348,	HEMBA1002381,	HEMBA1002458,	HEMBA1002462,
	HEMBA1002469,	HEMBA1002503,	HEMBA1002508.	HEMBA1002542,	HEMBA1002547,
	HEMBA1002552,	HEMBA1002558.	HEMBA1002583,	HEMBA1002590,	HEMBA1002592,
	HEMBA1002609,	HEMBA1002624,	HEMBA1002629.	HEMBA1002645,	HEMBA1002678,
45	HEMBA1002688,	HEMBA1002703,	HEMBA1002712,	HEMBA1002728,	HEMBA1002730,
45	HEMBA1002750,	HEMBA1002770,	HEMBA1002779,	HEMBA1002780,	HEMBA1002818.
	HEMBA1002833,	HEMBA1002876,	HEMBA1002934,	HEMBA1002935,	HEMBA1002968,
	HEMBA1002970,	HEMBA1002973, HEM	•	•	•
	HEMBA1003041,	HEMBA1003046,	HEMBA1003064.	HEMBA1003067,	HEMBA1003079,
50	HEMBA1003083,	HEMBA1003086,	HEMBA1003098,	HEMBA1003129,	HEMBA1003133,
	HEMBA1003142,	HEMBA1003166,	HEMBA1003175,	HEMBA1003179,	•
	HEMBA1003204.	HEMBA1003212,		HEMBA1003235,	HEMBA1003202,
	HEMBA1003286,	HEMBA1003212,	HEMBA1003220,	•	HEMBA1003257,
55	HEMBA1003280,	HEMBA1003296,	HEMBA1003322, HEMBA1003403,	HEMBA1003330,	HEMBA1003348,
	TEMENTAGOS (U,	UEMDY 1009910	TICMENTOUS 4US,	HEMBA1003418,	HEMBA1003447,

			LF 10/401/	7.6	
	HEMBA1003480,	HEMBA1003531,	HEMBA1003538, ·	HEMBA1003548,	HEMBA1003560,
	HEMBA1003571,	HEMBA1003591,	HEMBA1003597,	HEMBA1003615,	HEMBA1003621,
	HEMBA1003640,	HEMBA1003656,	HEMBA1003667,	HEMBA1003680,	HEMBA1003690,
5	HEMBA1003692,	HEMBA1003715,	HEMBA1003720,	HEMBA1003742,	HEMBA1003758,
	HEMBA1003803,	HEMBA1003805,	HEMBA1003827,	HEMBA1003836,	HEMBA1003838,
	HEMBA1003864,	HEMBA1003879,	HEMBA1003880,	HEMBA1003885,	HEMBA1003893,
	HEMBA1003902,	HEMBA1003926,	HEMBA1003937,	HEMBA1003939,	HEMBA1003950,
10	HEMBA1003958,	HEMBA1003989,	HEMBA1004000,	HEMBA1004012,	HEMBA1004015,
10	HEMBA1004024,	HEMBA1004048,	HEMBA1004056,	HEMBA1004074,	HEMBA1004111,
	HEMBA1004132,	HEMBA1004143.	HEMBA1004164,	HEMBA1004225,	HEMBA1004227,
	HEMBA1004238,	HEMBA1004246,	HEMBA1004267,	HEMBA1004274,	HEMBA1004289.
	HEMBA1004306,	HEMBA1004321,	HEMBA1004323,	НЕМВА1004335,	HEMBA1004353,
15	HEMBA1004354,	HEMBA1004356,	HEMBA1004408,	HEMBA1004429,	HEMBA1004499,
	HEMBA1004507,		· .	BA1004542, HEMBA1	•
	HEMBA1004586,	HEMBA1004596,	HEMBA1004604,	HEMBA1004629,	HEMBA1004669,
	HEMBA1004693,	HEMBA1004709,	HEMBA1004730,	HEMBA1004736,	HEMBA1004751,
20	HEMBA1004753,	HEMBA1004763,	HEMBA1004778,	HEMBA1004803,	HEMBA1004816,
	HEMBA1004850,	HEMBA1004880,	HEMBA1004923,	HEMBA1004930,	HEMBA1004954,
	HEMBA1004977,	HEMBA1004978,	HEMBA1004980,	HEMBA1005009,	HEMBA1005019,
	HEMBA1005035,	HEMBA1005079,	HEMBA1005123,	HEMBA1005133,	HEMBA1005149;
25	HEMBA1005202,	HEMBA1005206,	HEMBA1005219,	HEMBA1005241,	HEMBA1005296,
	HEMBA1005304,	HEMBA1005331,	HEMBA1005338,	HEMBA1005353,	HEMBA1005359,
	HEMBA1005374,	HEMBA1005382,	HEMBA1005403,	HEMBA1005408,	HEMBA1005411,
	HEMBA1005443,	HEMBA1005469,	HEMBA1005472,	HEMBA1005474,	HEMBA1005475,
30	HEMBA1005500,	HEMBA1005508,	HEMBA1005511,	HEMBA1005513,	HEMBA1005520,
	HEMBA1005526,	HEMBA1005528,	HEMBA1005548,	HEMBA1005552,	HEMBA1005568,
	HEMBA1005576,	HEMBA1005588,	HEMBA1005606,	HEMBA1005609,	HEMBA1005616,
	HEMBA1005627,	HEMBA1005631,	HEMBA1005634,	HEMBA1005670,	HEMBA1005679,
35	HEMBA1005680,	HEMBA1005705,	HEMBA1005765,	HEMBA1005813,	HEMBA1005822,
	HEMBA1005829,	HEMBA1005834,	HEMBA1005853,	HEMBA1005894,	HEMBA1005911,
	HEMBA1005921,	HEMBA1005931,	HEMBA1005934,	HEMBA1005990,	HEMBA1005991,
	HEMBA1005999,	HEMBA1006002,	HEMBA1006036,	HEMBA1006042,	HEMBA1006100,
40	HEMBA1006138,	HEMBA1006142,	HEMBA1006173,	HEMBA1006182,	HEMBA1006198,
70	HEMBA1006278,	HEMBA1006283, HE	MBA1006328, HEMI		*
	HEMBA1006377,	HEMBA1006380,	HEMBA1006381,	HEMBA1006416,	HEMBA1006419,
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45	HEMBA1006485,	HEMBA1006486,	HEMBA1006492,	HEMBA1006502,	HEMBA1006507,
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	PLACE1009230,	PLACE1009246,	PLACE1009298,	PLACE1009319,	PLACE1009328,
40	PLACE1009443,	PLACE1009468,	PLACE1009539,	PLACE1009595,	PLACE1009607,
	PLACE1009621,	PLACE1009622,	PLACE1009659,	PLACE1009763,	PLACE1009798,
	PLACE1009861,	PLACE1009879,	PLACE1009888,	PLACE1009924,	PLACE1009995,
	PLACE1009997,	PLACE1010023,	PLACE1010074,	PLACE1010076,	PLACE1010096,
45	PLACE1010105,	PLACE1010293,	PLACE1010310,	PLACE1010321,	PLACE1010329,
	PLACE1010383,	PLACE1010492,	PLACE1010522,	PLACE1010599,	PLACE1010616,
	PLACE1010624,	PLACE1010702,	PLACE1010720,	PLACE1010761,	PLACE1010771,
	PLACE1010786,	PLACE1010856, PL			
50	PLACE1010900,	PLACE1010925,	PLACE1010942,	PLACE1010947,	PLACE1010954,
•	PLACE1010960,	PLACE1010965,	PLACE1011026,	PLACE1011032,	PLACE1011054,
	PLACE1011056,	PLACE1011057,	PLACE1011090,	PLACE1011109,	PLACE1011114,
	PLACE1011185,	PLACE1011214,	PLACE1011291,	PLACE1011332,	PLACE1011340,
E E	PLACE1011371,	PLACE1011452,	PLACE1011477,	PLACE1011492,	PLACE1011567,
55	PLACE1011576,	PLACE1011586,	PLACE1011635,	PLACE1011646,	PLACE1011719,

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	PLACE1011729,	PLACE1011749,	PLACE1011783,	PLACE1011874,	PLACE1011923,
	PLACE1011962,	PLACE1011995,	PLACE2000003,	PLACE2000006,	PLACE2000011,
	PLACE2000014,	PLACE2000021,	PLACE2000030,	PLACE2000033,	PLACE2000034,
5	PLACE2000039,	PLACE2000047,	PLACE2000050,	PLACE2000062,	PLACE2000097,
~	PLACE2000100,	PLACE2000103,	PLACE2000111,	PLACE2000124,	PLACE2000140.
	PLACE2000170,	PLACE2000176,	PLACE2000187,	PLACE2000216,	PLACE2000235,
	PLACE2000246,	PLACE2000264,	PLACE2000302,	PLACE2000305,	PLACE2000335,
40	PLACE2000240,	PLACE2000342,	PLACE2000347,	PLACE2000366,	PLACE2000335,
10	PLACE2000341,	PLACE2000342,	PLACE2000347, PLACE2000399,	PLACE2000300,	PLACE2000371, PLACE2000411.
	PLACE2000419,	PLACE2000450,	PLACE2000458,	PLACE2000465,	PLACE3000004,
	PLACE3000009,	PLACE3000020,	PLACE3000029,	PLACE3000070,	PLACE3000103,
15	PLACE3000121,	PLACE3000124,	PLACE3000145,	PLACE3000147,	PLACE3000155,
	PLACE3000158,	PLACE3000160,	PLACE3000169,	PLACE3000194,	PLACE3000207,
	PLACE3000208,	PLACE3000220, PL	ACE3000221, PLAC	E3000226, PLACE3	000242,
	PLACE3000254,	PLACE3000271,	PLACE3000304,	PLACE3000320,	PLACE3000331,
20	PLACE3000339,	PLACE3000341,	PLACE3000350,	PLACE3000352,	PLACE3000353,
	PLACE3000362,	PLACE3000388,	PLACE3000399,	PLACE3000400,	PLACE3000401,
	PLACE3000405,	PLACE3000425,	PLACE3000455,	PLACE3000475,	PLACE3000477,
	PLACE4000009,	PLACE4000034,	PLACE4000049,	PLACE4000063,	PLACE4000089,
	PLACE4000100,	PLACE4000128,	PLACE4000131,	PLACE4000156,	PLACE4000211,
25	PLACE4000222,	PLACE4000230,	PLACE4000233,	PLACE4000250,	PLACE4000259,
	PLACE4000320,	PLACE4000323,	PLACE4000379,	PLACE4000411,	PLACE4000431,
	PLACE4000445,	PLACE4000450,	PLACE4000465,	PLACE4000487,	PLACE4000521,
	PLACE4000581,	SKNMC1000011,	SKNMC1000050,	SKNMC1000091,	THYR01000017,
30	THYR01000085,	THYRO1000092,	THYR01000132,	THYR01000163,	THYR01000186,
	THYR01000187,	THYR01000190,	THYR01000197,	THYR01000221,	THYR01000241,
	THYR01000253,	THYR01000320,	THYR01000394,	THYR01000438,	THYR01000452,
	THYR01000471,	THYR01000484,	THYR01000569,	THYR01000602,	THYR01000625,
35	THYR01000658,	THYR01000676,	THYR01000699,	THYR01000712,	THYR01000715,
	THYR01000815,	THYR01000829,	THYR01000843,	THYR01000855,	THYR01000865,
	THYR01000916,	THYR01000974,	THYR01000975,	THYR01000983,	THYR01000984,
	THYR01000988,	THYR01001003,	THYR01001031,	THYR01001093,	THYR01001133,
40	THYR01001173,	THYR01001177,	THYR01001189,	THYR01001204,	THYR01001213,
40	THYR01001287,	THYR01001320, TH	···		
	THYR01001405.	THYR01001406,	THYR01001411,	THYR01001426,	THYR01001480.
	THYRO1001487,	THYR01001534,	THYR01001537,	THYR01001541,	THYR01001559,
	THYR01001573,	THYR01001584,	THYR01001595,	THYR01001602,	THYR01001617,
45	THYR01001637,	THYR01001656,	THYR01001671,	THYR01001673,	THYR01001703,
	THYR01001706,	THYR01001738,	THYR01001772,	THYR01001793,	THYR01001828,
	THYR01001854,	THYR01001907,	VESEN10001122,	Y79AA1000033,	Y79AA1000037,
	Y79AA1000059,	Y79AA1000065,	Y79AA1000121,	Y79AA1000202,	Y79AA100037,
50	Y79AA1000039,	Y79AA1000065,	Y79AA1000131,	Y79AA1000342,	
	Y79AA1000231,	•		· ·	Y79AA1000346,
		Y79AA1000355,	Y79AA1000368,	Y79AA1000405,	Y79AA1000410,
	Y79AA1000420,	Y79AA1000469,	Y79AA1000480,	Y79AA1000538,	Y79AA1000539,
55	Y79AA1000560,	Y79AA1000589,	Y79AA1000627,	Y79AA1000705,	Y79AA1000734,
50	Y79AA1000774,	Y79AA1000782,	Y79AA1000784,	Y79AA1000827,	Y79AA1000833,

Y79AA1000850,	Y79AA1000966,	Y79AA1000968,	Y79AA1000985,	Y79AA1001061,
Y79AA1001068,	Y79AA1001077,	Y79AA1001078,	Y79AA1001105,	Y79AA1001145,
Y79AA1001211,	Y79AA1001216,	Y79AA1001228,	Y79AA1001236,	Y79AA1001299,
Y79AA1001394,	Y79AA1001402,	Y79AA1001511,	Y79AA1001533,	Y79AA1001548,
Y79AA1001555,	Y79AA1001581,	Y79AA1001585,	Y79AA1001603,	Y79AA1001647,
Y79AA1001665,	Y79AA1001679,	Y79AA1001711,	Y79AA1001805,	Y79AA1001827,
Y79AA1001846,	Y79AA1001866,	Y79AA1001875,	Y79AA1001923,	Y79AA1001963,
Y79AA1002089,	Y79AA1002093, Y7	9AA1002115, Y79A	A1002125, Y79AA1	002209,
Y79AA1002211,	Y79AA1002220,	Y79AA1002246,	Y79AA1002258,	Y79AA1002311,
Y79AA1002351,	Y79AA1002361, Y7	9AA1002472, Y79A	A1002482	

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Signal ratios of EC_AGE_BSA to EC_BSA and of EC_glycated_BSA to EC_BSA were calculated for each gene. Genes with high signal ratios were selected. In the case of calculating the ratio of signal value of 40 or less to that of more than 40, such signal values were, for convenience, taken as 40 instead of the real values. When the ratio EC_AGE_BSA/EC_BSA is 2 or more, expression of the genes exhibiting such ratio is expected to be elevated due to advanced glycation endproduct of bovine serum albumin. The higher the value is, the higher the gene expression level is. When the ratio EC_AGE_BSA/EC_BSA ranges from 0.5 to 2, expression of the genes exhibiting such ratio is expected to be unaffected due to advanced glycation endproduct of bovine serum albumin. When the ratio EC_AGE_BSA/EC_BSA is less than 0.5, expression of the genes exhibiting such ratio value is expected to be decreased due to advanced glycation endproduct of bovine serum albumin. The lower the value is, the lower the gene expression level is.

Clone with EC_AGE_BSA/EC_BSA ratio of 2 or higher are as follows: HEMBA1003958, MAMMA1001256, PLACE2000411.

Clone with EC_AGE_BSA/EC_BSA ratio of 0.5 or less is as follows: MAMMA1001783. These cDNAs are associated with diabetes.

When the ratio EC_glycated_BSA/EC_BSA is 2 or more, the expression level of the gene exhibiting such ratio is expected to be elevated due to glycated bovine serum albumin. The higher the value is, the higher the gene expression level is. When the ratio EC_glycated_BSA/EC_BSA ranges from 0.5 to 2, the expression level of the gene exhibiting such ratio is expected to be unaffected with glycated bovine serum albumin. When the ratio EC_glycated_BSA/EC_BSA is less than 0.5, the expression level of a gene exhibiting such ratio is expected to be decreased due to glycated bovine serum albumin. The lower the value is, the lower the gene expression level is.

Clones with EC_glycated_BSA/EC_BSA ratio of 2 or more are as follows: HEMBA1004850, MAMMA1001256, MAMMA1002132 and PLACE3000119.

A clone with EC_glycated_BSA/EC_BSA ratio of 0.5 or less is as follows: MAMMA1001783.

These cDNAs are also associated with diabetes.

Analysis of genes associated with neural cell differentiation

Genes involved in neural cell differentiation are useful for treating neurological diseases.

It is possible that genes with varying expression levels in response to induction of cellular differentiation in neural cells are associated with neurological diseases.

A survey was performed for genes of which expression levels are varied in response to induction of differentiation (stimulation by retinoic acid (RA)) in cultured cells of a neural strain, NT2.

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The NT2 cells were treated basically according to supplier's instruction manual. "Undifferentiated NT2 cells" means NT2 cells successively cultured in an Opti-MEM I (GIBCO-BRL; catalog No. 31985) containing 10%(v/v) fetal bovine serum and 1%(v/v) penicillinstreptomycin (GIBCO BRL). "NT2 cells cultured in the presence of retinoic acid" means the cells resulted from transferring undifferentiated NT2 cells into a retinoic acid-containing medium, which consists of D-MEM (GIBCO BRL; catalog No. 11965), 10%(v/v) fetal bovine serum, 1%(v/v) penicillin-streptomycin and 10 µM retinoic acid (GIBCO-BRL), and the subsequent successive culture therein for 5 weeks. "NT2 cells that were cultured in the presence of retinoic acid and then further cultured in the presence of cell-division inhibitor added" means NT2 cells resulted from transferring NT2 cells cultured in the presence of retinoic acid for 5 weeks into a celldivision inhibitor-containing medium, which consisted of D-MEM(GIBCO BRL; catalog No.11965), 10%(v/v) fetal bovine serum, 1%(v/v) penicillin-streptomycin, 10μM retinoic acid, 10 μM FudR (5fluoro-2' -deoxyuridine: GIBCO BRL), 10 µM Urd (Uridine: GIBCO BRL) and 1 µM araC (Cytosine β-D-Arabinofuranoside: GIBCO BRL), and the subsequence successive culture for 2 weeks. Each of the cells were treated with trypsin and then harvested. Total RNAs were extracted from the cells by using S.N.A.P. (TM) à Total RNA Isolation kit (Invitrogen (1) à). The labeling of probe used for hybridization was carried out by using 10 µg of the total RNA according to the same methods as described above. The data were obtained in triplicate (n=3). The data of signal value representing gene expression level in the cells in the presence of stimulation for inducing differentiation were compared with those in the absence of the stimulation. The comparison was performed by statistical treatment-of two-sample t-test. Clones with significant difference in the signal distribution were selected under the condition of p<0.05. In this analysis, clones with the difference can be statistically detected even when the signals were low. Accordingly, clones with signal value of 40 or less were also assessed for the selection.

Tables 170-349 show the expression level of each cDNA in undifferentiated NT2 cells, NT2 cells cultured in the presence of RA, and NT2 cells that were cultured in the presence of RA and that were further cultured in the presence of cell-division inhibitor added.

Averaged signal values (M_1, M_2) and sample variances $(s_1^2 \grave{a}, s_2^2 \grave{a})$ were calculated for each gene in each of the cells, and then, the pooled sample variances $s^2 \grave{a}$ were obtained from the sample variances of the two types of cells to be compared. The t values were determined according to the following formula: $t=(M1-M2)/s/(1/3+1/3)^{1/2} \grave{a}$. When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01 in the t-distribution table with 4 degrees of freeedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively. The tables also include the information on an increase (+) or decrease (-) in the expression level of a gene in the treated cells when the level is compared with that of untreated undifferentiated cells.

Clones of which expression levels increased by RA are as follows:
HEMBA1000005, HEMBA1000042, HEMBA1000046, HEMBA1000076, HEMBA1000111,
HEMBA1000141, HEMBA1000150, HEMBA1000185, HEMBA1000282, HEMBA1000304,

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	HEMBA1000307,	HEMBA1000338,	HEMBA1000357,	HEMBA1000376,	HEMBA1000387,
	HEMBA1000392,	HEMBA1000428,	HEMBA1000456,	HEMBA1000459,	HEMBA1000469,
	HEMBA1000504,	HEMBA1000508,	HEMBA1000519,	HEMBA1000540,	HEMBA1000545,
5	HEMBA1000557,	HEMBA1000563,	HEMBA1000568,	HEMBA1000575,	HEMBA1000588,
	HEMBA1000592,	HEMBA1000604,	HEMBA1000622,	HEMBA1000655,	HEMBA1000673,
	HEMBA1000682,	HEMBA1000726,	HEMBA1000727,	HEMBA1000749,	HEMBA1000769,
	HEMBA1000774,	HEMBA1000791,	HEMBA1000822,	HEMBA1000872,	HEMBA1000876,
10	HEMBA1000910,	HEMBA10009	•		BA1000960,
	HEMBA1000972,	HEMBA1000974,	HEMBA1000991	HEMBA1001008,	HEMBA1001020.
	HEMBA1001043,	HEMBA1001051,	HEMBA1001060,	HEMBA1001071,	HEMBA1001077
	HEMBA1001085,	HEMBA1001094,	HEMBA1001109,	HEMBA1001121,	HEMBA1001122.
15	HEMBA1001140.	HEMBA1001172,	HEMBA1001226,	HEMBA1001235,	HEMBA1001265,
	HEMBA1001281,	HEMBA1001294,	HEMBA1001299,	HEMBA1001319,	HEMBA1001323,
	HEMBA1001330,	HEMBA1001351,	HEMBA1001361,	HEMBA1001377,	HEMBA1001388,
	HEMBA1001391,	HEMBA1001398,	HEMBA1001432,	HEMBA1001435,	HEMBA1001442,
20	HEMBA1001454,	HEMBA1001455,	HEMBA1001497,	HEMBA1001517,	HEMBA1001569,
	HEMBA1001570,	HEMBA1001581,	HEMBA1001585,	HEMBA1001620,	HEMBA1001711,
	HEMBA1001718,	HEMBA1001723,	HEMBA1001761,	HEMBA1001815,	HEMBA1001819,
	HEMBA1001861,	HEMBA1001864,	HEMBA1001869,	HEMBA1001888,	HEMBA1001915,
25	HEMBA1001918,	HEMBA1001940,	HEMBA1001964,	HEMBA1001967,	HEMBA1001979,
	HEMBA1001987,	HEMBA1001991,	HEMBA1002008,	HEMBA1002022.	HEMBA1002039,
	HEMBA1002049,	HEMBA1002084,	HEMBA1002102,	HEMBA1002113,	HEMBA1002144,
	HEMBA1002160,	HEMBA1002162,	HEMBA1002185,	HEMBA1002212,	HEMBA1002226,
30	HEMBA1002229,	HEMBA1002267,	HEMBA1002270,	HEMBA1002337,	HEMBA1002381,
	HEMBA1002458,	HEMBA1002477,	HEMBA1002508,	HEMBA1002558,	HEMBA1002561,
	HEMBA1002583,	HEMBA1002590,	HEMBA1002628,	HEMBA1002645,	HEMBA1002661,
	HEMBA1002678,	HEMBA1002712,	HEMBA1002728,	HEMBA1002780,	HEMBA1002850,
35	HEMBA1002886,	HEMBA1002934,	HEMBA1002935,	HEMBA1002939,	HEMBA1002951,
	HEMBA1002968,	HEMBA1002970,	HEMBA1002973,	HEMBA1002999,	HEMBA1003021,
	HEMBA1003033,	HEMBA1003034,	HEMBA1003064,	HEMBA1003067,	HEMBA1003078,
	HEMBA1003086,	HEMBA1003096,	HEMBA1003129,	HEMBA1003142,	HEMBA1003148,
40	HEMBA1003166,	HEMBA1003175,	HEMBA1003197,	HEMBA1003199,	HEMBA1003202,
	HEMBA1003204,	HEMBA1003212,	HEMBA1003235,	HEMBA1003250,	HEMBA1003273,
	HEMBA1003276,	HEMBA1003278,	HEMBA1003291,	HEMBA1003309,	HEMBA1003322.
	HEMBA1003328,	HEMBA1003348,	HEMBA1003369,	HEMBA1003376,	HEMBA1003384,
45	HEMBA1003395,	HEMBA1003463,	HEMBA1003480,	HEMBA1003531,	HEMBA1003548,
	HEMBA1003591,	HEMBA1003595,	HEMBA1003597,	HEMBA1003617,	HEMBA1003621,
	HEMBA1003622,	HEMBA1003637,	HEMBA1003640,	HEMBA1003645,	HEMBA1003646,
	HEMBA1003656,	HEMBA1003679,	HEMBA1003692,	HEMBA1003715,	HEMBA1003720,
50	HEMBA1003725,	HEMBA1003729,	HEMBA1003758,	HEMBA1003803,	HEMBA1003805,
	HEMBA1003836,	HEMBA1003838,	HEMBA1003879,	HEMBA1003885,	HEMBA1003893,
	HEMBA1003908,	HEMBA1003937,	HEMBA1003942,	HEMBA1003953,	HEMBA1003958,
	HEMBA1003959,	HEMBA1003978,	HEMBA1003987,	HEMBA1003989,	HEMBA1004000,
55	HEMBA1004011,	HEMBA1004012,	HEMBA1004015,	HEMBA1004024,	HEMBA1004049,
	HEMBA1004056.	HEMBA1004111,	HEMBA1004132,	HEMBA1004143,	HEMBA1004164,
	HEMBA1004199,	HEMBA1004200,	HEMBA1004207,	HEMBA1004225.	HEMBA1004246.
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	HEMBA1004248,	HEMBA1004267,	HEMBA1004289,	HEMBA1004312,	HEMBA1004323,
	HEMBA1004335,	HEMBA1004353,	HEMBA1004354,	HEMBA1004356,	HEMBA1004366,
	HEMBA1004396,	HEMBA1004405,	HEMBA1004429,	HEMBA1004433,	HEMBA1004460,
5	HEMBA1004499,	HEMBA1004502,	HEMBA1004506,	HEMBA1004534,	HEMBA1004538,
	HEMBA1004573,	HEMBA1004577,	HEMBA1004586,	HEMBA1004610,	HEMBA1004629,
	HEMBA1004666,	HEMBA1004669,	HEMBA1004672,	HEMBA1004709,	HEMBA1004733,
	HEMBA1004736,	HEMBA1004748,	HEMBA1004751,	HEMBA1004758,	HEMBA1004763,
10	HEMBA1004768,	HEMBA1004770,	HEMBA1004778,	HEMBA1004803,	HEMBA1004820,
	HEMBA1004847,	HEMBA1004863,	HEMBA1004880,	HEMBA1004909,	HEMBA1004918,
	HEMBA1004923,	HEMBA1004930,	HEMBA1004934,	HEMBA1004944,	HEMBA1004954,
	HEMBA1004980,	HEMBA1005035,	HEMBA1005039,	HEMBA1005075,	HEMBA1005079,
15	HEMBA1005113,	HEMBA1005123,	HEMBA1005133,	HEMBA1005149,	HEMBA1005152,
	HEMBA1005219,	HEMBA1005223,	HEMBA1005232,	HEMBA1005251,	HEMBA1005274,
	HEMBA1005275,	HEMBA1005304,	HEMBA1005311,	HEMBA1005314,	HEMBA1005359,
	HEMBA1005367,	HEMBA1005374,	HEMBA1005410,	HEMBA1005411,	HEMBA1005423,
20	HEMBA1005426,	HEMBA1005443,	HEMBA1005447,	HEMBA1005474,	HEMBA1005508,
	HEMBA1005511,	HEMBA1005520,	HEMBA1005526,	HEMBA1005548,	HEMBA1005552,
	HEMBA1005576,	HEMBA1005581,	HEMBA1005582,	HEMBA1005583,	HEMBA1005588,
	HEMBA1005595,	HEMBA1005609,	HEMBA1005616,	HEMBA1005627,	HEMBA1005631,
25	HEMBA1005666,	HEMBA1005670,	HEMBA1005679,	HEMBA1005699,	HEMBA1005705,
	HEMBA1005765,	HEMBA1005780,	HEMBA1005822,	HEMBA1005829,	HEMBA1005834,
	HEMBA1005853,	HEMBA1005891,	HEMBA1005894,	HEMBA1005911,	HEMBA1005921,
	HEMBA1005991,	HEMBA1005999,	HEMBA1006036,	HEMBA1006042,	HEMBA1006100,
30	HEMBA1006138,	HEMBA1006142,	HEMBA1006268,	HEMBA1006278,	HEMBA1006328,
	HEMBA1006344,	HEMBA1006359,	HEMBA1006398,	HEMBA1006416,	HEMBA1006419,
	HEMBA1006421,	HEMBA1006426,	HEMBA1006438,	HEMBA1006483,	HEMBA1006485,
	HEMBA1006502,	HEMBA1006507,	HEMBA1006546,	HEMBA1006559,	HEMBA1006562,
35	HEMBA1006579,	HEMBA1006597,	HEMBA1006612,	HEMBA1006617,	HEMBA1006631,
	HEMBA1006635,	HEMBA1006652,	HEMBA1006695,	HEMBA1006744,	HEMBA1006754,
	HEMBA1006779,	HEMBA1006780,	HEMBA1006795,	HEMBA1006821,	HEMBA1006865,
	HEMBA1006926,	HEMBA1006973,	HEMBA1007017,	HEMBA1007052,	HEMBA1007080,
40	HEMBA1007085,	HEMBA1007113,	HEMBA1007121,	HEMBA1007147,	HEMBA1007149,
	HEMBA1007206,	HEMBA1007224,	HEMBA1007256,	HEMBA1007267,	HEMBA1007288,
	HEMBA1007327, I	HEMBA1007341, HE	MBA1007347,		
	HEMBB1000005,	HEMBB1000008,	HEMBB1000018,	HEMBB1000024,	HEMBB1000030,
45	HEMBB1000039,	HEMBB1000048,	HEMBB1000054,	HEMBB1000059,	HEMBB1000083,
	HEMBB1000089,	HEMBB1000099,	HEMBB1000113,	HEMBB1000141,	HEMBB1000144,
	HEMBB1000173,	HEMBB1000175,	HEMBB1000215,	HEMBB1000218,	HEMBB1000258,
	HEMBB1000264,	HEMBB1000272,	HEMBB1000307,	HEMBB1000338,	HEMBB1000339,
50	HEMBB1000343,	HEMBB1000354,	HEMBB1000374,	HEMBB1000376,	HEMBB1000420,
	HEMBB1000434,	HEMBB1000441,	HEMBB1000449,	HEMBB1000490,	HEMBB1000491,
	HEMBB1000523,	HEMBB1000530,	HEMBB1000550,	HEMBB1000554,	HEMBB1000573,
	HEMBB1000575,	HEMBB1000593,	HEMBB1000598,	HEMBB1000638,	HEMBB1000643,
55	HEMBB1000649,	HEMBB1000671,	HEMBB1000684,	HEMBB1000709,	HEMBB1000726,
	HEMBB1000770,	HEMBB1000774,	HEMBB1000790,	HEMBB1000835,	HEMBB1000840,
	HEMBB1000848,	HEMBB1000870,	HEMBB1000883,	HEMBB1000887,	HEMBB1000890,

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     HEMBB1001665,
                     MAMMA1000684,
                                     MAMMA1001139,
                                                      MAMMA1001743,
                                                                       NT2RM1000257,
     NT2RM1000318,
                     NT2RM1000539,
                                      NT2RM1000666,
                                                      NT2RM2000092,
                                                                       NT2RM2000192,
                                                                       NT2RM4001754,
     NT2RM2000371,
                     NT2RM2000594,
                                      NT2RM4000511,
                                                      NT2RM4001140,
     NT2RM4001905,
                     NT2RM4001940,
                                      NT2RM4002593,
                                                       NT2RP1000086,
                                                                       NT2RP1000439,
                     NT2RP2000098,
                                      NT2RP2000965,
                                                      NT2RP2001397,
                                                                       NT2RP2002047,
     NT2RP1001073,
                                                      NT2RP2005126,
                                                                       NT2RP2005464,
     NT2RP2004226.
                     NT2RP2004396.
                                      NT2RP2004655.
                                                      NT2RP3000980,
                                                                       NT2RP3001383,
     NT2RP2005712,
                     NT2RP2005859,
                                      NT2RP2005890,
     NT2RP3001621,
                     NT2RP3002081,
                                      NT2RP3002181,
                                                       NT2RP3002244,
                                                                       NT2RP3002590,
                     NT2RP3004258,
                                      NT2RP3004378,
                                                      NT2RP3004527,
                                                                       NT2RP3004594,
     NT2RP3003059,
                     NT2RP4001950,
                                      NT2RP4002047,
                                                       NT2RP4002408,
                                                                       NT2RP5003459,
     NT2RP4001760,
                     OVARC1000035,
                                      OVARC1000431,
                                                      OVARC1001051,
                                                                       OVARC1001129,
     OVARC1000004,
     OVARC1001176,
                     OVARC1001261,
                                      OVARC1001342,
                                                      OVARC1001942,
                                                                       OVARC1001943,
                                      PLACE1003190,
                                                      PLACE1003375,
                                                                       PLACE1004128,
     PLACE1002171,
                     PLACE1002465,
```

PLACE1005923,

PLACE1007257,

PLACE1007375,

PLACE1005026,

PLACE1005876,

PLACE1007507, PLACE1008941, PLACE1010624, PLACE1011090, PLACE1011219, THYRO1000270, Y79AA1000346, Y79AA1001541.

Clones of which expression levels increase in the presence of both RA and RA/inhibitor are as follows: HEMBA1000504. HEMBA1000588, HEMBA1000682, HEMBA1000046, HEMBA1000307, HEMBA1000726, HEMBA1000943, HEMBA1001071, HEMBA1001094, HEMBA1001122, HEMBA1001361, HEMBA1001455, HEMBA1001869, HEMBA1002084, HEMBA1001323. HEMBA1002583, HEMBA1002628, HEMBA1003096, HEMBA1003142, HEMBA1003276, HEMBA1003309. HEMBA1003463. HEMBA1003597. HEMBA1003617. HEMBA1003725, HEMBA1003803, HEMBA1003879, HEMBA1003989, HEMBA1004000, HEMBA1004015, HEMBA1004024, HEMBA1004049, HEMBA1004056, HEMBA1004199, HEMBA1004248, HEMBA1004356, HEMBA1004666, HEMBA1004770, HEMBA1004803, HEMBA1004923, HEMBA1004934, HEMBA1004954, HEMBA1005039, HEMBA1005075, HEMBA1005113, HEMBA1005367, HEMBA1005219, HEMBA1005232, HEMBA1005251, HEMBA1005304, HEMBA1005411, HEMBA1005548, HEMBA1005581, HEMBA1005631, HEMBA1005410, HEMBA1005666, HEMBA1005780, HEMBA1006485, HEMBA1006559, HEMBA1006579, HEMBB1000059, HEMBB1000575, HEMBB1000709, HEMBB1000848, HEMBA1006754, HEMBB1001210, HEMBB1001394, HEMBB1001668, HEMBB1000985, HEMBB1001117, HEMBB1001695, HEMBB1002049, HEMBB1002254, HEMBB1002266, HEMBB1002371, 25 HEMBB1002614, HEMBB1002617, HEMBB1002697, MAMMA1000241, MAMMA1000424, MAMMA1000616. MAMMA1000731, MAMMA1000824, MAMMA1001038, MAMMA1001243, MAMMA1002769, MAMMA1002871, MAMMA1002941, NT2RM1000725, NT2RM1000829, NT2RM2000504, NT2RM2000635, NT2RM2000821, 30 NT2RM2001582, NT2RM2001592, NT2RM2001632, NT2RM2001648, NT2RM2001671, NT2RM4000425, NT2RM4000471, NT2RM2001695, NT2RM2001879, NT2RM4000290, NT2RM4000531, NT2RM4000852, NT2RM4001047, NT2RM4001454, NT2RM4001557, NT2RM4001566, NT2RM4001582, NT2RM4001953, NT2RM4002409, NT2RM4002558, NT2RM4002594, NT2RP1000259, NT2RP1000418, NT2RP1000574, NT2RP1000782, NT2RP1000856, NT2RP1000943, NT2RP1000988. NT2RP1001013. NT2RP1001173, NT2RP1001546, NT2RP2000091, NT2RP2000208, NT2RP2000274. NT2RP2000329, NT2RP2001277. NT2RP2000369, NT2RP2000634, NT2RP2000987, NT2RP2001290, NT2RP2001366, NT2RP2001423, NT2RP2001436, NT2RP2001467, NT2RP2001506, NT2RP2001601, NT2RP2001926, NT2RP2001985, NT2RP2002041, NT2RP2002046, NT2RP2002078, NT2RP2002124, NT2RP2002193, NT2RP2002312, NT2RP2002316, NT2RP2002426, NT2RP2002457, NT2RP2002475, NT2RP2002520, NT2RP2002595, NT2RP2002701, NT2RP2003099, NT2RP2003137, NT2RP2003206, NT2RP2002672, NT2RP2003280, NT2RP2003393, NT2RP2003445, NT2RP2003230, NT2RP2003272, NT2RP2004681, NT2RP2003456, NT2RP2003596, NT2RP2003871, NT2RP2004743, NT2RP2004775, NT2RP2004933, NT2RP2004967, NT2RP2005003, NT2RP2005289, NT2RP2005453, NT2RP2005555. NT2RP2005767, NT2RP2005853, NT2RP2006043. NT2RP2006393, NT2RP2006436, NT2RP2006441, NT2RP2006467, NT2RP2006565, NT2RP3000359, NT2RP3000366, NT2RP3000403, NT2RP3000418, NT2RP3000441, NT2RP3000759, NT2RP3001007, NT2RP3001126, NT2RP3001355, NT2RP3000561, NT2RP3001396, NT2RP3001449, NT2RP3001490, NT2RP3001679, NT2RP3001752, NT2RP3001989, NT2RP3001782, NT2RP3001799, NT2RP3002142, NT2RP3002248, NT2RP3002343, NT2RP3002484, NT2RP3002529, NT2RP3002549, NT2RP3002687,

			E1 1 01 1 0 11 7 12		
	NT2RP3003032,	NT2RP3003139,	NT2RP3003193,	NT2RP3003204,	NT2RP3003210,
	NT2RP3003212,	NT2RP3003264,	NT2RP3003282,	NT2RP3003500,	NT2RP3004041,
	NT2RP3004215,	NT2RP4000147,	NT2RP4000259,	NT2RP4000360,	NT2RP4000448,
5	NT2RP4000524,	NT2RP400107	9, NT2RP40	01150, NT2R	P4001274,
	NT2RP4001353,	NT2RP4001547,	NT2RP4001677,	OVARC1000092,	OVARC1000321,
	OVARC1000384,	OVARC1000408,	OVARC1000414,	OVARC1000520,	OVARC1000526,
	OVARC1000588,	OVARC1000679,	OVARC1000682,	OVARC1000769,	OVARC1000862,
10	OVARC1000984,	OVARC1001000,	OVARC1001004,	OVARC1001154,	OVARC1001170,
•	OVARC1001173,	OVARC1001200,	OVARC1001268,	OVARC1001376,	OVARC1001476,
	OVARC1001542, 0	OVARC1001873, OV	ARC1001987, OVAI		,
	PLACE1000014,	PLACE1000048,	PLACE1000184,	PLACE1000246,	PLACE1000292,
15	PLACE1000332,	PLACE1000347,	PLACE1000712,	PLACE1001000,	PLACE1001294,
	PLACE1001311,	PLACE1001395,	PLACE1001570,	PLACE1001608,	PLACE1001610,
	PLACE1001746	PLACE1001821,	PLACE1001844,	PLACE1001897,	PLACE1002066,
	PLACE1002119,	PLACE1002157,	PLACE1002205,	PLACE1002256,	PLACE1002259,
20	PLACE1002438,	PLACE1002474,	PLACE1002477,	PLACE1002500,	PLACE1002578,
	PLACE1002815,	PLACE1002851,	PLACE1002968,	PLACE1003108,	PLACE1003174,
	PLACE1003200,	PLACE1003256,	PLACE1003334,	PLACE1003516,	PLACE1003592,
	PLACE1003723,	PLACE1003760,	PLACE1003771,	PLACE1003795,	PLACE1003892,
25	PLACE1003968,	PLACE1004103,	PLACE1004256,	PLACE1004629,	PLACE1004979,
	PLACE1005102,	PLACE1005128,	PLACE1005305,	PLACE1005477,	PLACE1005666,
	PLACE1005804,	PLACE1005934,	PLACE1006076,	PLACE1006119,	PLACE1006159,
	PLACE1006164,	PLACE1006170,	PLACE1006492,	PLACE1006629,	PLACE1006704,
30	PLACE1006760,	PLACE1006795,	PLACE1006962,	PLACE1007045,	PLACE1007386,
	PLACE1007416,	PLACE1007484,	PLACE1007544,	PLACE1007645,	PLACE1007743,
	PLACE1007807,	PLACE1007858,	PLACE1008002,	PLACE1008273,	PLACE1008368,
	PLACE1008532,	PLACE1008568,	PLACE1008696,	PLACE1008867,	PLACE1009027,
35	PLACE1009039,	PLACE1009298,	PLACE1009328,	PLACE1009621,	PLACE1009637,
	PLACE1010089,	PLACE1010106,	PLACE1010152,	PLACE1010491,	PLACE1010630,
	PLACE1010739,	PLACE1010896,	PLACE1010925,	PLACE1010965,	PLACE1011046,
	PLACE1011214,	PLACE1011433,	PLACE1011719,	PLACE1011762,	PLACE2000039,
40	PLACE2000216,	PLACE2000302,	PLACE2000342,	PLACE2000347,	PLACE2000379,
	PLACE3000121,	PLACE3000124,	PLACE3000242,	PLACE3000271,	PLACE3000362,
	PLACE3000365,	PLACE3000400,	PLACE3000401,	PLACE4000034,	PLACE4000089,
	PLACE4000522,	PLACE4000558,	THYRO1000197,	THYRO1000241,	THYRO1000327,
45	THYRO1000394,	THYRO1000488,	THYRO1000585,	THYRO1000625,	THYRO1000805,
	THYRO1000934,	THYRO1001133,	THYRO1001173,	THYRO1001213,	THYRO1001262,
	THYRO1001290,	Y79AA1000800,	Y79AA1000976,	Y79AA1001585,	Y79AA1001696,
	Y79AA1001711, Y	79AA1002258.			
50	Clones of which	h expression levels o	decrease in the pres	sence of both RA a	nd RA/inhibitor are
				HEMBA1005570,	NT2RM1000666,
	NT2RM2000092,	NT2RM2000594,	NT2RM4001754,		NT2RP2002047,
	NT2RP3000980,	NT2RP3002081,			
55		OVARC1001942,	·		
		PLACE1010624, Y79			
	•	*			

These are neurological disease-associated clones.

Analysis of rheumatoid arthritis-associated genes

5

The onset of rheumatoid arthritis is thought to be involved in the proliferation of synovial cells covering inner surfaces of joint cavity and in inflammatory reaction resulted from the action of cytokines produced by leukocytes infiltrating into the joint synovial tissues (Rheumatism Information Center, http://www.rheuma-net.or.jp/). Recent studies have also revealed that tissue necrosis factor (TNF)- α participates in the onset (Current opinion in immunology 1999; 11, 657-662). When the expression of a gene exhibits responsiveness to the action of TNF on synovial cells, the gene is considered to be involved in rheumatoid arthritis.

15

A survey was performed for genes of which expression levels are varied in response to TNF- α in the primary cell culture of synovial tissue. The primary cultured cells of the smooth muscle (Cell Applications) were grown to be confluent in a culture dish, and then, human TNF- α (Boehringer-Mannheim) was added at a final concentration of 10 ng/ml thereto. The culture was further continued for 24 hours.

20

Total RNA was extracted from the cells by using S.N.A.P. $^{(TM)}$ à Total RNA Isolation kit (Invitrogen). The labeling of probe used for hybridization was carried out by using 10 μ g of the total RNA according to the same methods as described above. The data were obtained in triplicate (n=3). The data of signal value representing gene expression level in the cells in the presence of TNF stimulation were compared with those in the absence of the stimulation. The comparison was performed by statistical treatment of two-sample t-test. Clones with significant difference in the signal distribution were selected under the condition of p<0.05. In this analysis,

30

25

clones with the difference can be statistically detected even when the signals were low. Accordingly, clones with signal value of 40 or less were also assessed for the selection.

Table 352 shows the expression level of each cDNA in synovial cells cultured in the absence or presence of TNF.

40

50

Averaged signal values (M_1, M_2) and sample variances $(s_1^{\ 2} \dot{a}, s_2^{\ 2} \dot{a})$ for each gene were calculated in each of the cells, and then, the pooled sample variances $s^2 \dot{a}$ were obtained from the sample variances of the two types of cells to be compared. The t-values were determined according to the following formula: $t=(M_1-M_2)/s/(1/3+1/3)^{1/2}\dot{a}$. When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01 in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively. The tables also include the information of an increase (+) or decrease (-) in the expression level of a gene in the stimulated cells when the level is compared with that of unstimulated cells.

Clones of which expression levels are elevated by TNF-\alpha are as follows:

HEMBA1000012, HEMBA1000020, HEMBA1000005, HEMBA1000046, HEMBA1000076, HEMBA1000111, HEMBA1000168, HEMBA1000185, HEMBA1000201, HEMBA1000231, HEMBA1000243, HEMBA1000280, HEMBA1000282, HEMBA1000304, HEMBA1000307, HEMBA1000327, HEMBA1000356, HEMBA1000376, HEMBA1000387, HEMBA1000390, HEMBA1000418, HEMBA1000460, HEMBA1000491, HEMBA1000501, HEMBA1000518, HEMBA1000519. HEMBA1000520, HEMBA1000531. HEMBA1000534. HEMBA1000542. HEMBA1000545, HEMBA1000591, HEMBA1000592, HEMBA1000594, HEMBA1000636,

			E. 10,4011 /12		
	HEMBA1000655,	HEMBA1000657,	HEMBA1000673,	HEMBA1000682,	HEMBA1000686,
	HEMBA1000722,	HEMBA1000726,	HEMBA1000827,	HEMBA1000870,	HEMBA1000918,
	HEMBA1000971,	HEMBA1000974,	HEMBA1000986,	HEMBA1001019,	HEMBA1001043,
5	HEMBA1001051,	HEMBA1001059,	HEMBA1001060,	HEMBA1001071,	HEMBA1001080,
	HEMBA1001109,	HEMBA1001140,	HEMBA1001172,	HEMBA1001196,	HEMBA1001213,
	HEMBA1001226,	HEMBA1001281,	HEMBA1001299,	HEMBA1001302,	HEMBA1001303,
	HEMBA1001323,	HEMBA1001326,	HEMBA1001327,	HEMBA1001330,	HEMBA1001351,
10	HEMBA1001407,	HEMBA1001411,	HEMBA1001446,	HEMBA1001454,	HEMBA1001569,
	HEMBA1001647,	HEMBA1001714,	HEMBA1001800,	HEMBA1001804,	HEMBA1001809,
	HEMBA1001888,	HEMBA10019			BA1001967,
	HEMBA1002084,	HEMBA1002161,	HEMBA1002166,	HEMBA1002241,	HEMBA1002337,
15	HEMBA1002363,	HEMBA1002389,	HEMBA1002458,	HEMBA1002460,	HEMBA1002469,
	HEMBA1002538,	HEMBA1002542,	HEMBA1002547,	HEMBA1002609,	HEMBA1002624,
	HEMBA1002659,	HEMBA1002750,	HEMBA1002770,	HEMBA1002779,	HEMBA1002810,
	HEMBA1002816,	HEMBA1002818,	HEMBA1002850,	HEMBA1002863,	HEMBA1003021,
20	HEMBA1003033,	HEMBA1003078,	HEMBA1003166,	HEMBA1003202,	HEMBA1003204,
	HEMBA1003229,	HEMBA1003235,	HEMBA1003276,	HEMBA1003286,	HEMBA1003296,
	HEMBA1003370,	HEMBA1003376,	HEMBA1003403,	HEMBA1003418,	HEMBA1003433,
	HEMBA1003447,	HEMBA1003560,	HEMBA1003569.	HEMBA1003571,	HEMBA1003591,
25	HEMBA1003597,	HEMBA1003598,	HEMBA1003621,	HEMBA1003656,	HEMBA1003662,
	HEMBA1003680,	HEMBA1003715,	HEMBA1003725,	HEMBA1003729,	HEMBA1003733,
	HEMBA1003742,	HEMBA1003773,	HEMBA1003783,	HEMBA1003950,	HEMBA1004012,
	HEMBA1004015,	HEMBA1004048,	HEMBA1004074,	HEMBA1004086,	HEMBA1004111,
30	HEMBA1004131,	HEMBA1004202,	HEMBA1004203,	HEMBA1004207,	HEMBA1004248,
	HEMBA1004274,	HEMBA1004321,	HEMBA1004330,	HEMBA1004356,	HEMBA1004366,
	HEMBA1004405,	HEMBA1004408,	HEMBA1004429,	HEMBA1004499,	HEMBA1004507,
	HEMBA1004509,	HEMBA1004542,	HEMBA1004596,	HEMBA1004604,	HEMBA1004776,
35	HEMBA1004889,	HEMBA1004934,	HEMBA1004978,	HEMBA1005019,	HEMBA1005047,
	HEMBA1005206,	HEMBA1005219,	HEMBA1005274,	HEMBA1005331,	HEMBA1005338,
	HEMBA1005394,	HEMBA1005423,	HEMBA1005576,	HEMBA1005732,	HEMBA1005746,
	HEMBA1006091,	HEMBA1006142,	HEMBA1006173,	HEMBA1006198,	HEMBA1006253,
40	HEMBA1006268,	HEMBA1006309,	HEMBA1006377,	HEMBA1006474,	HEMBA1006486,
,-	HEMBA1006492,	HEMBA1006502,	HEMBA1006535.	HEMBA1006579,	HEMBA1006648.
	HEMBA1006659,	HEMBA1006885,	HEMBA1006929,	HEMBA1006941,	HEMBA1007078,
			HEMBA1007194.		
45	HEMBA1007080,	HEMBA1007121,		HEMBA1007300,	HEMBA1007301,
	HEMBA1007322,	HEMBB1000036,	HEMBB1000044,	HEMBB1000089,	HEMBB1000215,
	HEMBB1000217,	HEMBB10002/2,	HEMBB1000420,	HEMBB1000591,	HEMBB1000593,
	HEMBB1000631,	HEMBB1000835,	HEMBB1000887,	HEMBB1000908,	HEMBB1000975,
50	HEMBB1000985,	HEMBB1001011,	HEMBB1001014,	HEMBB1001112,	HEMBB1001133,
	HEMBB1001331,	HEMBB1001337,	HEMBB1001366,	HEMBB1001367,	HEMBB1001384,
	HEMBB1001394,	HEMBB1001429,	HEMBB1001463,	HEMBB1001619,	HEMBB1001684,
	HEMBB1001706,	HEMBB1001753,	HEMBB1001797,	HEMBB1001802,	HEMBB1001812,
55	HEMBB1001874,	HEMBB1001910,	HEMBB1001915,	HEMBB1001973,	HEMBB1001983,
	HEMBB1001990,	HEMBB1002190,	HEMBB1002193,	HEMBB1002249,	HEMBB1002329,
	HEMBB1002342,	HEMBB1002371,	HEMBB1002409,	HEMBB1002442,	HEMBB1002489,
	HEMBB1002492,	HEMBB1002520,	HEMBB1002534,	HEMBB1002596,	HEMBB1002664,

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	HEMBB1002692,	HEMBB1002697,	HEMBB1002705,	MAMMA1000092,	MAMMA1000155,
	MAMMA1000163,	MAMMA1000173,	MAMMA1000175,	MAMMA1000227,	MAMMA1000241,
	MAMMA1000257,	MAMMA1000264,	MAMMA1000266,	MAMMA1000270,	MAMMA1000307,
5	MAMMA1000410,	MAMMA1000413,	MAMMA1000416,	MAMMA1000421,	MAMMA1000472,
	MAMMA1000501,	MAMMA1000605,	MAMMA1000643,	MAMMA1000670,	MAMMA1000684,
	MAMMA1000696,	MAMMA1000732;	MAMMA1000752,	MAMMA1000802,	MAMMA1000824,
	MAMMA1000905,	MAMMA1000921,	MAMMA1000931,	MAMMA1000957,	MAMMA1000962,
10	MAMMA1000998,	MAMMA1001008,	MAMMA1001050,	MAMMA1001074,	MAMMA1001078,
	MAMMA1001292,	MAMMA1001397,	MAMMA1001476,	MAMMA1001743,	MAMMA1001744,
	MAMMA1001754,	MAMMA1001760,	MAMMA1001785,	MAMMA1001858,	MAMMA1001908,
	MAMMA1002236,	MAMMA1002267,	MAMMA1002292,	MAMMA1002311,	MAMMA1002322,
15	MAMMA1002359,	MAMMA1002362,	MAMMA1002485,	MAMMA1002494,	MAMMA1002597,
	MAMMA1002598,	MAMMA1002665,	MAMMA1002671,	MAMMA1002684,	MAMMA1002748,
	MAMMA1002775,	MAMMA1002830,	MAMMA1002858,	MAMMA1002868,	MAMMA1002886,
	MAMMA1002887,	MAMMA1002892,	MAMMA1002909,	MAMMA1002937,	MAMMA1002947,
20	MAMMA1002964,	MAMMA1002970,	MAMMA1003013,	MAMMA1003150,	NT2RM1000039,
	NT2RM1000062,	NT2RM1000080,	NT2RM1000086,	NT2RM1000127,	NT2RM1000132,
	NT2RM1000187,	NT2RM1000199,	NT2RM1000244,	NT2RM1000256,	NT2RM1000272,
	NT2RM1000318,	NT2RM1000354,	NT2RM1000377,	NT2RM1000430,	NT2RM1000499,
25	NT2RM1000539,	NT2RM1000553,	NT2RM1000563,	NT2RM1000699,	NT2RM1000742,
	NT2RM1000826,	NT2RM1000829,	NT2RM1000833,	NT2RM1000882,	NT2RM1000898,
	NT2RM1000905,	NT2RM1001092,	NT2RM2000013,	NT2RM2000032,	NT2RM2000042,
	NT2RM2000101,	NT2RM2000124,	NT2RM2000192,	NT2RM2000259,	NT2RM2000260,
30	NT2RM2000363,	NT2RM2000368,	NT2RM2000402,	NT2RM2000452,	NT2RM2000952,
	NT2RM2001221,	NT2RM2002014,	NT2RM2002030,	NT2RM4000156,	NT2RM4000349,
	NT2RM4000395,	NT2RM4000457,	NT2RM4000511,	NT2RM4000514,	NT2RM4000698,
	NT2RM4000764,	NT2RM4001016,	NT2RM4001084,	NT2RM4001594,	NT2RM4001629,
35	NT2RM4001662,	NT2RM4001841,	NT2RM4002093,	NT2RM4002109,	NT2RM4002145,
	NT2RM4002189,	NT2RM4002194,	NT2RM4002226,	NT2RP1000170,	NT2RP1000439,
	NT2RP1000478,	NT2RP1000513,	NT2RP1000701,	NT2RP1000856,	NT2RP1001361,
40	NT2RP2000097,	NT2RP2000239,	NT2RP2000288,	NT2RP2000328,	NT2RP2000329,
40	NT2RP2000369,	NT2RP2000422,	NT2RP2000842,	NT2RP2000965,	NT2RP2001245,
	NT2RP2001440,	NT2RP2001560,	NT2RP2001634,	NT2RP2001663,	NT2RP2001677,
	NT2RP2001762,	NT2RP2002270,	NT2RP2002312,	NT2RP2002316,	NT2RP2002333,
45	NT2RP2002706,	NT2RP2002925,	NT2RP2002959,	NT2RP2002987,	NT2RP2003125,
70	NT2RP2003137,	N12RP2003237,	NT2RP2003272,	NT2RP2003596,	NT2RP2003604,
	N12RP2003643,	N12RP2003968,	NT2RP2003976,	NT2RP2004194,	NT2RP2004321,
	NT2RP2005037,	NT2RP2005140,	N12RP2005204,	NT2RP2005293,	NT2RP2005457,
50	NT2RP2005555,	NT2RP2005600,	NT2RP2005701,	NT2RP2005719,	NT2RP2005722,
	NT2RP2005773,	NT2RP2005890,	NT2RP2006023,	NT2RP2006071,	NT2RP3000186,
	NT2RP3000341,	NT2RP3000599,	NT2RP3000632,	NT2RP3000644,	NT2RP3000852,
	NT2RP3000968,	NT2RP3001096,	NT2RP3001109,	NT2RP3001126,	NT2RP3001147,
55	NT2RP3001449, NT2RP3001969,	NT2RP3001529, NT2RP3002081,	NT2RP3001753, NT2RP3002142,	NT2RP3001854,	NT2RP3001915,
	NT2RP3001909, NT2RP3002603,	NT2RP3002081, NT2RP3002810,	NT2RP3002142, NT2RP3002876,	NT2RP3002399,	NT2RP3002590,
	NT2RP3002603, NT2RP3003672,	NT2RP3002810, NT2RP3004209,	NT2RP3002878,	NT2RP3003311,	NT2RP3003330,
	1412NF3003012,	N12NF3004209,	1412RF3004378,	NT2RP4000078,	NT2RP4000541,

			LI I UI T UII ME		
	NT2RP4000588,	NT2RP4001219,	NT2RP4001228,	NT2RP4001276,	NT2RP4001507,
	NT2RP4002047,	NT2RP5003459,	NT2RP5003492,	OVARC1000085,	OVARC1000087,
	OVARC1000106,	OVARC1000151,	OVARC1000198,	OVARC1000431,	OVARC1000440,
5	OVARC1000564,	OVARC1000605,	OVARC1000679,	OVARC1000883,	OVARC1000912,
	OVARC1000960,	OVARC1000971,	OVARC1001038,	OVARC1001055,	OVARC1001085,
	OVARC1001129,	OVARC1001167,	OVARC1001339,	OVARC1001425,	OVARC1001745,
	OVARC1001762,	OVARC1001766,	OVARC1001942,	OVARC1002044,	OVARC1002138,
10	PLACE1000004,	PLACE1000005,	PLACE1000420,	PLACE1000547,	PLACE1000562,
•	PLACE1000653,	PLACE1001168,	PLACE1001311,	PLACE1001377,	PLACE1001920,
	PLACE1001983,	PLACE1002066,	PLACE1002072,	PLACE1002140,	PLACE1002171,
	PLACE1002319,	PLACE1002474,	PLACE1002499,	PLACE1002532,	PLACE1002665,
15	PLACE1003025,	PLACE1003145,	PLACE1003361,	PLACE1003605,	PLACE1003704,
	PLACE1003783,	PLACE1003885,	PLACE1004405,	PLACE1004629,	PLACE1004686,
	PLACE1004930,	PLACE1005066,	PLACE1006077,	PLACE1005630,	PLACE1005876,
	PLACE1006143,	PLACE1006325,	PLACE1006488,	PLACE1006805,	PLACE1006829,
20	PLACE1007286,	PLACE1007858,	PLACE1008201,	PLACE1009045,	PLACE1009113,
	PLACE1009621,	PLACE1010106,	PLACE1010310,	PLACE1010622,	PLACE1010944,
	PLACE1010965,	PLACE1011185,	PLACE1011332,	PLACE1011635,	PLACE1011646,
	PLACE1011725,	PLACE2000014,	PLACE2000264,	PLACE2000394,	PLACE2000419,
25	PLACE3000160,	PLACE3000220,	PLACE3000254,	PLACE3000271,	PLACE3000339,
	PLACE3000341,	PLACE3000350,	PLACE3000353,	PLACE3000401,	PLACE4000300,
	SKNMC1000091,	THYRO1000855,	THYRO1001559,	Y79AA1000065,	Y79AA1000202,
	Y79AA1000214,	Y79AA1000346,	Y79AA1000784,	Y79AA1000833,	Y79AA1000968,
30	Y79AA1001555, Y7	79AA1002220.			
	Clones of which	expression levels	decrease by TNF- $lpha$ a	are as follows:	
	HEMBA1002150,	HEMBB1000240,	NT2RM2000469, I	NT2R <mark>M</mark> 2000984, N	T2RM2001688, _
	NT2RM4000290,	NT2RM4000496,	NT2RM4000590,	NT2RM4001047,	NT2RM4001582,
35	NT2RM4001611,	NT2RM4001650,	NT2RM4002075,	NT2RM4002128,	NT2RP1000174,
	NT2RP1000243,	NT2RP1000581,	NT2RP1000688,	NT2RP1000767,	NT2RP1000825,
	NT2RP1001185,	NT2RP1001286,	NT2RP1001432,	NT2RP1001457,	NT2RP2000001,
	NT2RP2000248,	NT2RP2000841,	NT2RP2001813,	NT2RP2002137,	NT2RP2002928,
40	NT2RP2003517,	NT2RP2003559,	NT2RP2003564,	NT2RP2004933,	NT2RP2005038,
	NT2RP2006365,	NT2RP3000072,	NT2RP3000320,	NT2RP3000484,	NT2RP3000980,
	NT2RP3001111,	NT2RP3001420,	NT2RP3001495,	NT2RP3002056,	NT2RP3002057,

NT2RP3002713, NT2RP3002799, NT2RP3002869, NT2RP3002953, NT2RP3002545, 45 NT2RP3002955, NT2RP3003282, NT2RP3003290, NT2RP3003384, NT2RP3003385, NT2RP4000500, NT2RP3003870, NT2RP3004207, NT2RP3004262, NT2RP3004527, NT2RP4000524, NT2RP4000787, NT2RP4000927, NT2RP4000955, NT2RP4000989, NT2RP4001442, NT2RP4001638, NT2RP4001950, NT2RP4002888, NT2RP5003524,

50 OVARC1001270, PLACE1000246, PLACE1002816.

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These are rheumatoid arthritis-associated clones.

Analysis of ultraviolet radiation damage-associated genes

It is known that ultraviolet rays give considerably adverse influence on the health. In

recent years, there have been significant risks of tissue damage by ultraviolet rays because of destruction of the ozone layer. Thus, ultraviolet radiation has been recognized as a risk factor for skin diseases such as skin cancers (United States Environmental Protection Agency: Ozone Depletion Home Page, http://www.epa.gov/ozone/). Genes of which expression levels are varied in skin epidermal cells exposed to ultraviolet rays are considered to be associated with skin damage caused by ultraviolet radiation.

After primary cultured skin fibroblast cells were irradiated with ultraviolet ray and were cultured, a survey was performed for genes of which expression levels were varied depending on the irradiation of ultraviolet ray. First, after cultured to be confluent, the primary cultured skin fibroblast cells (Cell Applications) were exposed to 10,000 μ J/cm²à of 254-nm ultraviolet light.

Messenger RNAs were, then, extracted by using a FastTrack™ 2.0 mRNA Isolation kit (Invitrogen Co.) from the unexposed cells and from the cells that were exposed to the ultraviolet light and then cultured for 4 or 24 hours. The labeling of the hybridization probe was carried out by using a 1.5 µg of each mRNA in the same manner as described above. The data were obtained in triplicate (n=3). The hybridization signals were compared between the cells exposed to the ultraviolet light and the unexposed cells. The comparison was preformed by statistical treatment with two-sample t-test. Clones with significant differences in the signal distribution were selected under the condition of p<0.05. In this analysis, even when the signal is lower than others, the difference in the signal values can be detected statistically. Accordingly, clones with signal value of 40 or lower were also assessed for selection.

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Tables 353-509 show the expression of each cDNA in skin-derived fibroblast cells exposed and unexposed to ultraviolet light.

Averaged signal values (M_1, M_2) and sample variances $(s_1^{\ 2}\dot{a}, s_2^{\ 2}\dot{a})$ were calculated for each gene in each of the cells, and then, the pooled sample variances $s^2\dot{a}$ were obtained from the sample variances of the two types of cells to be compared. The t values were determined according to the following formula: $t=(M_1-M_2)/s/(1/3+1/3)^{1/2}\dot{a}$. When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01 in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively. The tables also include the information of an increase (+) or decrease (-) in the expression level of a gene in the exposed cells in comparison with that of unexposed cells.

The expression levels of the following clones were elevated 4 or 24 hours after the ultraviolet irradiation:

HEMBA1000542, HEMBA1002177, HEMBA1001808, HEMBA1003314, MAMMA1001874. NT2RM2001100, NT2RP2005732, NT2RP3000592, NT2RP4000657, OVARC1000004, NT2RM4001002, NT2RM4001813, OVARC1001092, OVARC1001342, PLACE1002816, NT2RM4002266, NT2RP2001174, NT2RP2001196, NT2RP2005358, NT2RP3000690, NT2RP3001216, NT2RP3003464, PLACE1006382, THYRO1000070, THYRO1001100, Y79AA1000342.

The expression levels of the following clones were decreased 4 or 24 hours after the ultraviolet irradiation:

HEMBA100005, HEMBA1000150, HEMBA1000156, HEMBA1000158, HEMBA1000168, HEMBA1000301, HEMBA1000304, HEMBA1000307, HEMBA1000333, HEMBA1000366, HEMBA1000369, HEMBA1000390, HEMBA1000396, HEMBA1000418, HEMBA1000434,

			E		
	HEMBA1000464,	HEMBA1000469,	HEMBA1000490,	HEMBA1000504,	HEMBA1000505,
	HEMBA1000557,	HEMBA1000657,	HEMBA1000673,	HEMBA1000682,	HEMBA1000686,
	HEMBA1000727,	HEMBA1000752,	HEMBA1000851,	HEMBA1000852,	HEMBA1000870,
5	HEMBA1000872,	HEMBA1001085,	HEMBA1001121,	HEMBA1001133,	HEMBA1001235,
	HEMBA1001265,	HEMBA1001281,	HEMBA1001289,	HEMBA1001299,	HEMBA1001303,
	HEMBA1001310,	HEMBA1001323,	HEMBA1001595,	HEMBA1001620,	HEMBA1001640,
	HEMBA1001678,	HEMBA1001712,	HEMBA1001835,	HEMBA1001950,	HEMBA1001987,
10	HEMBA1002253,	HEMBA1002321,	HEMBA1002341,	HEMBA1002419,	HEMBA1002679,
	HEMBA1002728,	HEMBA1002818,	HEMBA1002935,	HEMBA1002999,	HEMBA1003034,
	HEMBA1003071,	HEMBA1003098,	HEMBA1003142,	HEMBA1003175,	HEMBA1003202,
	HEMBA1003212,	HEMBA1003220,	HEMBA1003276,	HEMBA1003373,	HEMBA1003417,
15	HEMBA1003447,	HEMBA1003528,	HEMBA1003684,	HEMBA1003799,	HEMBA1003885,
	HEMBA1003989,	HEMBA1004011,	HEMBA1004055,	HEMBA1004133,	HEMBA1004225,
	HEMBA1004272,	HEMBA1004353,	HEMBA1004631,	HEMBA1004669,	HEMBA1004705,
	HEMBA1004753,	HEMBA1004776,	HEMBA1004803,	HEMBA1004816,	HEMBA1004900,
20	HEMBA1005047,	HEMBA10050	79, HEMBA10	005101 HEMI	BA1005149,
	HEMBA1005152,	HEMBA1005202,	HEMBA1005314,	HEMBA1005372,	HEMBA1005511,
	HEMBA1005513,	HEMBA1005518,	HEMBA1005570,	HEMBA1005577,	HEMBA1005581,
	HEMBA1005588,	HEMBA1005609,	HEMBA1005632,	HEMBA1005853,	HEMBA1006031,
25	HEMBA1006035,	HEMBA1006485,	HEMBA1006486,	HEMBA1006502,	HEMBA1006696,
	HEMBA1006789,	HEMBA1006796,	HEMBA1007085,	HEMBA1007224,	HEMBA1007301,
	HEMBA1007319,	HEMBA1007341,	HEMBA1007342,	HEMBB1000036,	HEMBB1000037,
	HEMBB1000217,	HEMBB1000266,	HEMBB1000317,	HEMBB1000336,	HEMBB1000354,
30	HEMBB1000369,	HEMBB1000399,	HEMBB1000434,	HEMBB1000438,	HEMBB1000592,
	HEMBB1000673,	HEMBB1000789,	HEMBB1000810,	HEMBB1000883,	HEMBB1000887,
	HEMBB1001105,	HEMBB1001182,	HEMBB1001242,	HEMBB1001267,	HEMBB1001424,
	HEMBB1001464,	HEMBB1001531,	HEMBB1001618,	HEMBB1001996,	HEMBB1002092,
35	HEMBB1002139,	HEMBB1002142,	HEMBB1002190,	HEMBB1002453,	HEMBB1002520,
	HEMBB1002550,	HEMBB1002556,	HEMBB1002600,	HEMBB1002664,	MAMMA1000009,
	MAMMA1000055,	MAMMA1000069,	MAMMA1000133,	MAMMA1000171,	MAMMA1000173,
	MAMMA1000287,	MAMMA1000416,	MAMMA1000585,	MAMMA1000713,	MAMMA1000760,
40	MAMMA1000798,	MAMMA1000831,	MAMMA1000875,	MAMMA1000876,	MAMMA1000877,
	MAMMA1000906,	MAMMA1000931,	MAMMA1000962,	MAMMA1001133,	MAMMA1001139,
	MAMMA1001243,	MAMMA1001271,	MAMMA1001274,	MAMMA1001298,	MAMMA1001606,
	MAMMA1001630,	MAMMA1001670,	MAMMA1001743,	MAMMA1001751,	MAMMA1002140,
45	MAMMA1002145,	MAMMA1002158,	MAMMA1002170,	MAMMA1002236,	MAMMA1002311,
	MAMMA1002498,	MAMMA1002754,	MAMMA1002780,	MAMMA1002820,	MAMMA1002843,
	MAMMA1002844,	MAMMA1002871,	MAMMA1003047,	NT2RM1000037,	NT2RM1000039,
	NT2RM1000080,	NT2RM1000086,	NT2RM1000341,	NT2RM1000499,	NT2RM1000669,
50	NT2RM1000746,	NT2RM1000781,	NT2RM1000885,	NT2RM1000905,	NT2RM1000962,
	NT2RM2000239,	NT2RM2000260,	NT2RM2000371,	NT2RM2000639,	NT2RM2000649,
	NT2RM2000735,	NT2RM2000821,	NT2RM2000984,	NT2RM2001035,	NT2RM2001065,
66	NT2RM2001105,	NT2RM2001177,	NT2RM2001194,	NT2RM2001196,	NT2RM2001243,
55	NT2RM2001256,	NT2RM2001424,	NT2RM2001588,	NT2RM2001635,	NT2RM2001648,
	NT2RM2001652,	NT2RM2001668,	NT2RM2001706,	NT2RM2001727,	NT2RM2001730,
	NT2RM2001743,	NT2RM2001753,	NT2RM2001760,	NT2RM2001771,	NT2RM2001785,

	NT2RM2001800,	NT2RM2001855,	NT2RM2001896,	NT2RM2001997,	NT2RM2002030,
	NT2RM2002049,	NT2RM2002091,	NT2RM2002142,	NT2RM2002145,	NT2RM2002178,
	NT2RM2002580,	NT2RM4000215,	NT2RM4000344,	NT2RM4000368,	NT2RM4000421,
5	NT2RM4000425,	NT2RM4000457,	NT2RM4000496,	NT2RM4000515,	NT2RM4000712,
	NT2RM4000787,	NT2RM4000813,	NT2RM4000820,	NT2RM4000852,	NT2RM4000950,
	NT2RM4000996,	NT2RM4001016,	NT2RM4001047,	NT2RM4001054,	NT2RM4001140,
	NT2RM4001151,	NT2RM4001187,	NT2RM4001204,	NT2RM4001258,	NT2RM4001437,
10	NT2RM4001454,	NT2RM4001489,	NT2RM4001605,	NT2RM4001611,	NT2RM4001666,
	NT2RM4001710,	NT2RM4001714,	NT2RM4001715,	NT2RM4001731,	NT2RM4001741,
	NT2RM4001746,	NT2RM4001856,	NT2RM4001938,	NT2RM4001940,	NT2RM4001984,
	NT2RM4001987,	NT2RM4002013,	NT2RM4002054,	NT2RM4002055,	NT2RM4002073,
15	NT2RM4002145,	NT2RM4002146,	NT2RM4002161,	NT2RM4002174,	NT2RM4002194,
	NT2RM4002205,	NT2RM4002213,	NT2RM4002266,	NT2RM4002278,	NT2RM4002558,
	NT2RM4002565,	NT2RM4002593,	NT2RM4002594,	NT2RP1000063,	NT2RP1000111,
	NT2RP1000124,	NT2RP1000174,	NT2RP1000348,	NT2RP1000443,	NT2RP1000522,
20	NT2RP1000677,	NT2RP1000695,	NT2RP1000730,	NT2RP1000767,	NT2RP1000834,
	NT2RP1000856,	NT2RP1000944,	NT2RP1001014,	NT2RP1001073,	NT2RP1001079,
	NT2RP1001177,	NT2RP1001185,	NT2RP1001199,	NT2RP1001253,	NT2RP1001310,
	NT2RP1001313,	NT2RP1001385,	NT2RP1001410,	NT2RP1001449,	NT2RP1001546,
25	NT2RP2000120,	NT2RP2000126,	NT2RP2000205,	NT2RP2000224,	NT2RP2000274,
	NT2RP2000298,	NT2RP2000310,	NT2RP2000327,	NT2RP2000328,	NT2RP2000329,
	NT2RP2000337,	NT2RP2000369,	NT2RP2000414,	NT2RP2000498,	NT2RP2000510,
	NT2RP2000715,	NT2RP2000819,	NT2RP2000931,	NT2RP2000938,	NT2RP2001044,
30	NT2RP2001065,	NT2RP2001081,	NT2RP2001137,	NT2RP2001312,	NT2RP2001327,
	NT2RP2001366,	NT2RP2001381,	NT2RP2001420,	NT2RP2001427,	NT2RP2001449,
	NT2RP2001511,	NT2RP2001526,	NT2RP2001560,	NT2RP2001576,	NT2RP2001581,
	NT2RP2001601,	NT2RP2001613,	NT2RP2001628,	NT2RP2001663,	NT2RP2001675,
35	NT2RP2001677,	NT2RP2001678,	NT2RP2001699,	NT2RP2001721,	NT2RP2001740,
	NT2RP2001813,	NT2RP2001861,	NT2RP2001876,	NT2RP2001907,	NT2RP2001926,
	NT2RP2001946,	NT2RP2001985,	NT2RP2002032,	NT2RP2002046,	NT2RP2002076,
	NT2RP2002099,	NT2RP2002105,	NT2RP2002137,	NT2RP2002172,	NT2RP2002208,
40	NT2RP2002219,	NT2RP2002231,	NT2RP2002256,	NT2RP2002270,	NT2RP2002292,
	NT2RP2002316,	NT2RP2002333,	NT2RP2002520,	NT2RP2002549,	NT2RP2002621,
	NT2RP2002706,	NT2RP2002710,	NT2RP2002750,	NT2RP2002800,	NT2RP2002862,
	NT2RP2002880,	NT2RP2002925,	NT2RP2002929,	NT2RP2002939,	NT2RP2002959,
45	NT2RP2002993,	NT2RP2003000,	NT2RP2003158,	NT2RP2003243,	NT2RP2003265,
	NT2RP2003277,	NT2RP2003280,	NT2RP2003295,	NT2RP2003297,	NT2RP2003433,
	NT2RP2003445,	NT2RP2003456,	NT2RP2003506,	NT2RP2003543,	NT2RP2003559,
	NT2RP2003567,	NT2RP2003604,	NT2RP2003706,	NT2RP2003777,	NT2RP2003781,
50	NT2RP2003871,	NT2RP2003952,	NT2RP2003976,	NT2RP2004013,	NT2RP2004014,
	NT2RP2004081,	NT2RP2004142,	NT2RP2004165,	NT2RP2004170,	NT2RP2004194,
	NT2RP2004196,	NT2RP2004239,	NT2RP2004240,	NT2RP2004321,	NT2RP2004364,
E E	NT2RP2004365,	NT2RP2004373,	NT2RP2004389,	NT2RP2004392,	NT2RP2004399,
55	NT2RP2004538,	NT2RP2004587,	NT2RP2004600,	NT2RP2004602,	NT2RP2004614,
	NT2RP2004664,	NT2RP2004675,	NT2RP2004681,	NT2RP2004710,	NT2RP2004743,
	NT2RP2004767,	NT2RP2004799,	NT2RP2004816,	NT2RP2004959,	NT2RP2004962,

	NT2RP2005000,	NT2RP2005003,	NT2RP2005012,	NT2RP2005031,	NT2RP2005037,
	NT2RP2005108,	NT2RP2005126,	NT2RP2005144,	NT2RP2005147,	NT2RP2005289,
	NT2RP2005293,	NT2RP2005325,	NT2RP2005354,	NT2RP2005441,	NT2RP2005464,
5	NT2RP2005491,	NT2RP2005495,	NT2RP2005498,	NT2RP2005501,	NT2RP2005555,
	NT2RP2005557,	NT2RP2005605,	NT2RP2005622,	NT2RP2005645,	NT2RP2005651,
	NT2RP2005654,	NT2RP2005669,	NT2RP2005694,	NT2RP2005701,	NT2RP2005752,
	NT2RP2005773,	NT2RP2005804,	NT2RP2005835,	NT2RP2005841,	NT2RP2005857,
10	NT2RP2005890,	NT2RP2005901,	NT2RP2005942,	NT2RP2006023,	NT2RP2006052,
	NT2RP2006069,	NT2RP2006166,	NT2RP2006184,	NT2RP2006186,	NT2RP2006196,
	NT2RP2006238,	NT2RP2006258,	NT2RP2006312,	NT2RP2006320,	NT2RP2006565,
	NT2RP2006571,	NT2RP3000050,	NT2RP3000055,	NT2RP3000072,	NT2RP3000080,
15	NT2RP3000109,	NT2RP3000134,	NT2RP3000142,	NT2RP3000149,	NT2RP3000186,
	NT2RP3000197,	NT2RP3000207,	NT2RP3000220,	NT2RP3000247,	NT2RP3000251,
	NT2RP3000361,	NT2RP3000366,	NT2RP3000418,	NT2RP3000449,	NT2RP3000451,
	NT2RP3000484,	NT2RP3000487,	NT2RP3000512,	NT2RP3000527,	NT2RP3000542,
20	NT2RP3000561,	NT2RP3000562,	NT2RP3000578,	NT2RP3000596,	NT2RP3000605,
	NT2RP3000624,	NT2RP3000644,	NT2RP3000661,	NT2RP3000665,	NT2RP3000739,
	NT2RP3000850,	NT2RP3000868,	NT2RP3000980,	NT2RP3001084,	NT2RP3001096,
	NT2RP3001109,	NT2RP3001111,	NT2RP3001115,	NT2RP3001119,	NT2RP3001120,
25	NT2RP3001126,	NT2RP3001140,	NT2RP3001155,	NT2RP3001176,	NT2RP3001272,
	NT2RP3001297,	NT2RP3001325,	NT2RP3001339,	NT2RP3001398,	NT2RP3001447,
	NT2RP3001457,	NT2RP3001459,	NT2RP3001495,	NT2RP3001554,	NT2RP3001580,
	NT2RP3001587,	NT2RP3001608,	NT2RP3001672,	NT2RP3001676,	NT2RP3001712,
30	NT2RP3001730,	NT2RP3001782,	NT2RP3001799,	NT2RP3001929,	NT2RP3001943,
	NT2RP3002002,	NT2RP3002004,	NT2RP3002045,	NT2RP3002054,	NT2RP3002057,
	NT2RP3002142,	NT2RP3002163,	NT2RP3002165,	NT2RP3002173,	NT2RP3002273,
	NT2RP3002351,	NT2RP3002631,	NT2RP3002659,	NT2RP3002671,	NT2RP3002713,
35	NT2RP3002955,	NT2RP3003071,	NT2RP3003078,	NT2RP3003139,	NT2RP3003242,
	NT2RP3003301,	NT2RP3003302,	NT2RP3003313,	NT2RP3003330,	NT2RP3003377,
	NT2RP3003384,	NT2RP3003411,	NT2RP3003701,	NT2RP3003831,	NT2RP3003870,
	NT2RP3003992,	NT2RP3004013,	NT2RP3004070,	NT2RP3004148,	NT2RP3004206,
40	NT2RP3004207,	NT2RP3004215,	NT2RP3004332,	NT2RP3004341,	NT2RP3004349,
	NT2RP3004399,	NT2RP4000023,	NT2RP4000051,	NT2RP4000129,	NT2RP4000147,
	NT2RP4000212,	NT2RP4000214,	NT2RP4000323,	NT2RP4000449,	NT2RP4000515,
	NT2RP4000519,	NT2RP4000528,	NT2RP4000556,	NT2RP4000588,	NT2RP4000638,
45	NT2RP4000648,	NT2RP4000973,	NT2RP4000996,	NT2RP4001006,	NT2RP4001029,
	NT2RP4001041,	NT2RP4001100,	NT2RP4001117,	NT2RP4001174,	NT2RP4001235,
	NT2RP4001274,	NT2RP4001276,	NT2RP4001315,	NT2RP4001373,	NT2RP4001389,
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50	NT2RP4001677,	NT2RP4001753,	NT2RP4001927,	NT2RP4002052,	NT2RP4002058,
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10	PLACE1009338,	PLACE1009375,	PLACE1009434,	PLACE1009613,	PLACE1009637,
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	THYRO1000197,	THYRO1000221,	THYRO1000253,	THYRO1000270,	· THYRO1000279,
	THYRO1000327,	THYRO1000394,	THYRO1000438,	THYRO1000558,	THYRO1000569,
35	THYRO1000585,	THYRO1000596,	THYRO1000625,	THYRO1000637,	THYRO1000676,
	THYRO1000734,	THYRO1000777,	THYRO1000783,	THYRO1000805,	THYRO1000843,
	THYRO1000934,	THYRO1001033,	THYRO1001347,	THYRO1001405,	THYRO1001411,
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	Y79AA1001541,	Y79AA1001585,	Y79AA1001705,	Y79AA1001781,	Y79AA1001923,
15	1/70444004000 1/		* * * * * * * * * * * * * * * * * * *		407

⁴⁵ Y79AA1001963, Y79AA1002125, Y79AA1002229, Y79AA1002407, Y79AA1002487.

These clones are ultraviolet radiation damage-associated clones.

Table 12

Expression of each cDNA in human tissues (containing clones that are not described in Examples.)

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	Clone name	None 1	lune (P. gland	Thymus	Brain	Kidney	Liver	Spleen
	GAPOH (Cr1)	38. 210	32.670	23.820	13. 580	11, 230	21, 120	24.910	22.440
	Bactin(Cr2)	279. 280	368.870		117. 500	92.880	114.650	82.990	256.790
	ADRGL 1000005	53.882	23.005	32.749	22.858	26.564	24.940	22.644	27.001
5	ADRGL 1000007	94.778	85. 185	160. 457	67. 191	101.768	62.489	67.150	73.543
3	ADRGL 1000009	11.141	50. 520-	10. 357	7.177	6.013	5. 219	14. 272	21.225
	ADRGL1000011	71.656	24. 579	29. 358	19.473	24.898	30.747	49. 220	22. 221
	ADRGL1000017	36.238	25. 252	20. 855	7.328	11.196	14, 298	19.658	11.288
	ADRGL1000058	66.209	129. 497	55. 226	49. 241	30.219	55.872	67.027	243. 436
	ADRGL1000069	38.630	23. 459	28. 991	12.540	27.353	33.633	28.774	20.911
10	ADRGL1000077	97.465	63.656	448. 427	83.412	71.108	53.740	67.906	89.439
	ADRGL 1000092	89. 423	45. 692	55. 810	26.033	44, 148	73.339	96.037	73.091
	ADRGL 1000099	73.675	24. 424	36.128	17.024	25. 964	41. 391	42.837	29.666
	ADRGL 1000136	141.745	63.974	77.017	24.777	33.549	58. 986	295, 009	84. 985
	ADRGL 1000147	394.563	155.829	271.210	92.899	165. 627	251.266	253. 420	150. 294
45	ADRGL 1000159	50.073	25.425	39. 296	15.194	16.125	20.040	33.720	23. 278
15	ADRGL 1000160	69.386	31.051	59.416	20.154	39.799	27.027	47. 169	20.716
	ADRGL1000171	57.047	23.011	43.063	23.860	40.581	59.814	117.055	32.630
	ADRGL1000181	45. 892	18.666	34. 476	15.434	34. 225	32. 962	39.693	16. 334
	BGG111000015	153. 242	42.337	92.865	41.003	45. 168	88. 524	85. 990	73.392
	BGG111000016	177.367	94.731	119. 688	34.159	30.249	98.805	98.783	39. 204
20	BGG111000017	84.712	32.614	38. 131	20.878	18.769	32.340	39.666	20.750
	BGG111000022	52.468	20.452	67.167	12.167	11.158	18.241	19.197	11.937
	BGG111000031	30.008	17.072	40.883	12.585	13.313	15, 525	16.757	13.406
	BGG111000042	49. 926	36.336	51.176	25.964	43. 122	43.770	49. 107	38.776
	BGG111000046	31.618	26.472	34. 182	31.854	12.650	25.784	18. 430	25. 385
	BNCH41000020	5031, 103	2993.496	1444. 841	537.162	6973.542		3350. 527.	3649. 144
25	BNGH41000025	91.717	35.026	73. 901	27.713	30.765	36.523	37.596	47.074
	BNGH41000026	176, 757	77.439	98. 345	35. 807	56.991	91.310	75.797	70.241 23.859
	BNGH41000027	65. 029	56. 353	25. 896	22.494	12.763	23.748	17.836	64.458
	BNCH41000035	148,779	66.776	119. 727	56.576	60.995	96.959 36.272	72.461	24.771
	BNGH41000037	79. 500	29.611	43. 438	18.317	20.857 102.259	125.323	86.783	122.959
30	BNGH41000042	224. 484	110.084	168. 448	14.729	27. 312	22.435	29.566	28. 937
••	BNG141000048	56. 144	32. 253	54.063 30.075	15.602	10.072	20, 735	16, 100	7.642
	BNGH41000056	67. 258	18.694 46.173	77.657	35.329	40.900	50.029	50.841	45. 285
	BNGH41000087	98. 262	16.985	28. 392	10.147	5.469	22.794	10.725	12.410
	BNGH41000091	50.895 69.043	34.730	40. 597	18.088	27.072	22.074	25.410	24.950
	BNGH41000157 BNGH41000169	44, 850	21.770	28. 655	11.403	25.991	28.509	25.634	25. 843
35	BNGH41000181	17, 163	15. 689	13. 948	3.996	9. 287	13.139	15. 553	16. 575
	BNGH41000198	81. 510	36.250	60.860	20.585	26.929	35.751	31.695	28. 325
	BNCH41000219	30. 302	25. 156	22. 187	13.757	11.208	15. 235	27.285	35.709
	BNGH41000229	252.790	65, 948	93. 499	51, 108	92.555	101.245	96.716	78. 256
	BNGH41000237	85.757	46, 997	55. 170	26.780	33.764	47.456	37.007	39.131
40	BNGH41000238	17.744	36.938	42.350	14.922	35.749	42.848	39. 238	13. 241
,0	BNGH41000243	45. 446	23.667	44.798	20.875	10.516	23.918	22. 443	27.033
	BNGH41000270	60, 889	18.651	29.618	10.724	15.979	12.351	19. 152	22.314
	BRAWH1000004	43.673	28. 539	7.640	11.388	19.198	14. 903	32.353	23.777
	BRAWH1000018	59.409	17. 941	102. 270	17, 107	709.078	25.732	24. 214	24.767
	BRAWH1000021	104.772	29, 951	51.142	21.042	1169.154	55.762	66.754	27.969
45	BRAWH1000027	152, 205	47.310	67.089	32. 199	64. 521	70.731	79.670	40. 928 100. 132
	BRAWH1000029	106. 376	49, 221	55. 840	40.856	59.552	56.487	17.998	15. 196
	BRAWH1000040	29. 419	16.761	31. 101	16.622	30.633	18. 200 97. 720	81, 271	69. 194
	BRAWH1000050	161. 264	71.786	118. 976	51.863 20.726	30.434	42.055	53. 856	24.624
	BRAWH1000051	74.067	34. 341	44. 047 35. 012		19.127	38.662		28. 094
50	8RAWH100006D	68. 789			8. 901	18.133	17.219		
30	BRAWH1000075	17.318	16.898 12.998				17.673		
	BRAWH1000081	43.025					77.111	110.167	
	BRAWH1000084 BRAWH1000095	174. 384					53.814		
	BRAWH1000095	146. 112					52.880		
	BRAWH1000097						73.707		
55	BRAWH1000100						16.582		
	BRAWH1000101	134. 838					71.642		
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	BRAWH1000104 BRAWH1000107	25.414 16.949	18. 303 5. 616	14.825	7.695 5.518	6. 355	5. 084	9, 107	6. 573
	BRA#H1000110	615, 476	492.704		383,512	368.156	369.621	277.348	340, 450
5	BRAWH1000111	175, 556	68. 459	92.209	45.974	64.703	81.723	90.369	57.301
•	BRAWH1000135	199.303	38, 098	72.093	26.809	57.720	91.668	87.016	35.866
	BRAWH1000190	56.386	41.640	57.914	22.782	55.671	40.034	35. 280	40. 134
	HEMBA1000005	11.985	23. 427	18.882	9.766	12.656	9. 959	23.443	21.677
	HEMBA1000006	37. 398	24, 521	24. 529	15, 587	22.317	13.336	16.038	15. 295
	HEMBA1000012	81.820	57.193	66.828	26.683	55. 423	58.731	85.614	66.259
10	HEMBA1000020	157.967	64. 157	115.635	51.940	77.293	77. 321	83.989	74. 362
	HEMBA1000030	82.882	35. 447	66.058	26.464	40. 990	60.871	47.058	50.652
	HEMBA1000034	47.434	17.878	50.696	5.594	14.005	6.673	24.652	7.134
	HEMBA 1 000042	147.376	94.003	330.908	69.071	76.472	55.477	37.783	60.479
	HEMBA 1 000045	28.478	20. 289	20.548	12, 445	11.835	22.788	11.196	15.775
15	HEMBA1000046	85.160	84. 475	242.940	57.017	68. 488	45. 288	37.098	47.486
10	HEMBA1000047	21.380	18.899	18.166	11.393	11. 185	12. 292	6.491	12.018 37.362
	HEMBA1000048	243.559	55.114	84.448	24. 247	43. 131	99. 333	57.041 10.347	7.770
	HEMBA1000050	22.711	11.878	21.972	7.477	4. 096 30. 754	13. 675 36. 740	34, 184	24. 269
	HEMBA1000053	45.071	26.410 ₄ 34.766	38.158 50.643	15.982 19.938	34, 641	54. 061	42.309	22. 530
	HEMBA1000060 HEMBA1000072	101. 197 240. 166	213.938	224.688	163.030	115. 246	207. 809	112.361	276.098
20	HEMBA1000073	23, 202	9. 580	10.815	1.698	6, 680	18.155	12.304	14. 973
	HEMBA1000076	95. 997	46. 783	177.931	32.617	48.964	50.792	33.947	44.142
	HEMBA1000084	66.603	25.710	48. 434	18.005	22. 553	38.118	40.479	29.683
	HENBA1000087	70.084	17.515	26.544	8.450	17.590	29. 220	19.519	22.565
	HEMBA1000088	15.474	8.614	19.903	4,775	4.519	11.446	34.905	6. 528
25	HEMBA1000091	80.622	38.604	59. 393	23.956	44. 939	49.760	33.946	24.614
25	HEMBA1000111	85.814	95. 270	270.642	75.147	54. 384	70.071	29. 529	55. 422
	HEMBA1000121	55.476	43, 368	146.465	37.419	29. 198	30.694	17.702	30. 398 23. 965
	HEMBA1000128	37.278	27.165 19.659	34.516 44.907	13.519	17.702	28.069 30.959	24. 383	26. 851
	HEMBA1000129	12, 961	24. 515	32.107	14.353	13.502	11.152	8. 907	20.635
	HEMBA1000141 HEMBA1000146	29, 273	11.479	20.418	8. 202	9.575	14.877	10.000	7.817
30	HEMBA1000150	534. 562	326. 814	684, 147	211.774	218.448	322.246	235.752	256.883
	HEMBA1000154	95.272	92, 253	101.483	54. 276	42.896	75. 526	92.689	188.019
	HEMBA1000156	50.177	72.591	58.026	31.149	21.865	38. 964	27.634	50.220
	HEMBA1000158	260.718	63.920	89.680	36.337	44. 915	93.421	111.344	53. 562
	HEMBA1000168	74.416	61.152	62.826	30.512	23. 287	34.966	44.005	33.564
35	HEMBA 1000180	28.502	22. 412	28.571	11.701	19, 230	10.903	11.731	14, 102
33	HEMBA 1000185	115.723	50.661	213.994	51.166	43. 435	56. 261	38.862	44. 992
	HEMBA1000188	21. 302	14. 879	16.948	11.392	11.821	10.656 3.349	12.501 8.544	7.842
	HEMBA1000193	14. 122	8. 318 49. 534	11.905	7.519	4.736	34, 328	23.359	56. 497
	HEMBA1000194 HEMBA1000201	54.688 21.062	14. 098	8.690	6.237	3, 109	5.059	9.317	10.522
	HEMBA1000211	22.388	25. 532	25.777	8.470	17.320	9.084	8, 469	11.766
40	HEMBA1000216	65.935	51.368	92.680	19.202	33.659	40. 971	36. 328	34.891
	HEMBA1000227	52.577	31. 332	34.925	19.503	18.411	21.504	22,590	25. 781
	HEMBA1000231	114.369	54. 299	131.256	38.550	43.246	29.778	24. 266	30.410
	HEMBA1000237	91.024	91, 360	199.338	58. 292	93. 250	57.000	49, 319	59. 288
	HEMBA1000243	53.456	43. 969	117.519	38, 431	25. 396	32.604	38, 910	32. 153
45	HEMBA1000244	173.469	104, 733	115. 584	33.079	65.527	124, 532	90. 927	78.610
43	HEMBA1000251	22.709	12. 333	14.367	9.019	16.095	13. 221	11.516 20.553	11.018
	HEMBA1000254	74.060	35. 626	130.009	20.848	37. 481 3. 863	24, 002	13, 690	3.797
	HEMBATO00264	29.478	15. 248	23. 537	9.473	9, 705	12. 833	7.348	24. 793
	HEMBA1000269	36.718	13. 465 39. 367	84.077	38.845	77.871	49. 267	36.211	38.871
	HEMBA1000275 HEMBA1000280	66.201 33.299	36. 073	54. 357	24.720	38.017	35.751	21.696	30.785
50	HEMBA1000282	93.815	121. 083	171.037	93.484	123.971	70.384	56.916	92.414
	HEMBA1000287	12.439	24. 935	29.793	10.840	37. 925	9.632	2.866	7.311
	HEMBA 1 000288	45. 269	30.009	145. 363	25.471	9.769	16.272	9.701	15.510
	HEMBA1000290	14.803	5.750	10.615	5.725	2.559	8. 602	8. 358	9.224
	HEMBA 1000298	27.085	22. 525	21.195	9.790	16.909	12.402	15. 289	17.159
<i></i>	HEMBA 1000300	98. 491	119, 119	304.884	73.660	85. 595	48, 175	43. 496	66.547
55	HEMBA1000302	23.840	15. 442	27.722	16.143	13.081	13.879	8. 259	12.569

REBARLO00310 129, 986 51, 013 88, 777 97, 211 504, 492 42, 618 43, 517 57, 77 57, 78 57,								00 004 1	44 010 1	40 271
FREERITORISTY COSK 22,013 31,964 13,157 15,571 7,974 10,014 8,685 FREERITORISTY 97,002 69,310 83,922 45,322 45,027 53,007 52,968 37,741 38,266 FREERITORISTY 75,605 55,618 63,147 41,731 53,007 43,526 69,421 46,566 FREERITORISTY 75,605 55,618 63,147 41,731 53,004 48,526 69,421 46,566 FREERITORISTY 75,605 75,518 53,148 53,004 48,526 69,421 46,566 FREERITORISTY 75,605 75,772 346,775 55,468 34,911 43,666 34,666 76,666		HEMBA 1000303	129. 286	51.013	88.777	32. 513	50.462	82.994	44.818	49. 271
FREEDRINGS 18 18 18 18 18 18 18 1										
	_	HEMBA1000307	14.054							
FEBRITODIST 19	5	HEMBA1000312	97.082	69.330	183.923					
		HEMBA1000318	16. 164	16.264	18.766	11.688				
FEBRAT000339 121 678 62 577 344,75 55. 463 49 114 38. 561 39. 598 40. 544 FEBRAT000349 27. 661 72. 585 31.755 7. 920 17. 555 11. 522 14. 407 15. 611 FEBRAT000349 27. 661 72. 585 31.755 7. 920 17. 555 11. 522 14. 407 15. 611 FEBRAT000349 27. 661 72. 585 31.755 7. 920 17. 555 11. 522 14. 407 15. 611 FEBRAT000359 57. 678 57. 578 57. 74 17. 572 37. 598 37. 578 37. 598 37. 5		HEMBA1000327	29.404	59.618	81.347	41,731				
FEBRAT000338 12, 878 62, 572 348, 751 55, 463 49, 114 38, 561 30, 598 40, 644		HENBA1000333	16.964	13.930	14.530	1. 872	5.776			
FEBRAT000143				62.572	348, 751	55, 463	49.114	38. 561	30.698	40.644
							5.872	16. 551	10.139	14.088
	10						17.658	11. 522-	14. 807	15.611
Figuration 15 15 15 15 15 15 15 1								35. 370	27.645	28, 615
FEBRAT0003157 84.8 81.7 86.2 42.7 20.784 17.5 81.4 31.0 88.4 44.431 28.8 89.9								43.511	37.808	26.628
									40. 431	28. 899
									26, 076	
HEBBA10003159									14,632	23, 547
	15								37, 622	36, 900
HEBBA1000318										
20 HEBRATOROSS 100.487 129.367, 181.196 20.287 100.485 14.007 57.619 79.876 HEBRATOROSS 59.342 34.021 71.118 22.346 27.319 47.916 53.026 34.161 HEBRATOROSS 19.400 22.884 44.179 8.776 11.742 10.594 12.865 12.631 HEBRATOROSS 19.400 22.884 44.179 8.776 11.742 10.594 12.265 12.631 HEBRATOROSS 19.400 22.884 44.179 8.776 11.742 10.594 12.265 12.631 HEBRATOROSS 19.400 59.81 870 27.799 30.933 11.216 50.056 18.580 HEBRATOROSS 18.65 10.778 14.987 4.031 12.495 7.918 15.386 17.771 19.844 HEBRATOROSS 19.400 18.500 18.200 19.800 19.800 19.800 19.800 19.800 19.800 19.800 19.800 HEBRATOROSS 19.400 18.500 18.500 18.500 18.500 18.500 HEBRATOROSS 19.400 19.800 18.200 19.800 19.800 19.800 19.800 19.800 HEBRATOROSS 19.400 19.800 19										
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HEIBA1000542 64.656 33.152 55.093 30.174 35.278 55.508 47.917 47.623 HEIBA1000545 148.870 136.401 48.802 8.499 12.534 7.119 25.484 15.094 HEIBA1000547 14.825 20.199 32.694 7.058 22.359 12.020 13.535 20.227 HEIBA1000551 163.806 171.089 543.876 131.764 115.775 116.646 69.596 152.516 HEIBA1000555 10.531 20.199 25.801 24.488 14.071 15.431 5.986 10.933 HEIBA1000557 80.051 48.396 168.724 37.150 32.863 31.872 22.800 30.926								17.606		
HEMBA1000542 64.656 33.152 58.093 30.174 35.278 55.508 47.917 47.623 HEMBA1000545 148.870 136.401 48.802 8.499 12.534 7.119 25.484 15.094 HEMBA1000547 14.825 20.199 32.694 7.058 22.359 12.020 13.535 20.227 HEMBA1000551 163.806 171.089 543.876 131.764 115.775 116.646 69.596 152.516 HEMBA1000555 10.531 20.199 25.801 24.488 14.071 15.431 5.986 10.933 168.724 37.150 32.863 31.872 22.800 30.926 32.272 33.272 3	50	HEMBA 1000540								
HEMBA1000547 14.825 20.199 32.694 7.058 22.359 12.020 13.535 20.227 HEMBA1000551 163.806 171.089 543.876 131.764 115.775 116.646 69.596 152.516 HEMBA1000555 10.531 20.199 25.801 24.488 14.071 15.431 5.986 10.933 168.724 37.150 32.863 31.872 22.800 30.926	-		64.656	33. 152						
HEMBA1000547			148.870							
HEMBA1000551 163.806 171.089 543.876 131.764 115.775 116.646 69.596 152.516 HEMBA1000555 10.531 20.199 25.801 24.488 14.071 15.431 5.986 10.933 HEMBA1000557 80.051 48.396 168.724 37.150 32.863 31.872 22.800 30.926				20. 199						
HEMBA1000555 10.531 20.199 25.801 24.488 14.071 15.431 5.986 10.933 HEMBA1000557 80.051 48.396 168.724 37.150 32.863 31.872 22.800 30.926										
55 HEMBA1000557 80.051 48.396 168.724 37.150 32.863 31.872 22.800 30.926					25.801					
	55			48.396	168.724					
	- -				51.047	10. 187	16.301	34.904	24.661	22.470
		<u> </u>								

	HEMBA1000563	9.473	11.545	18. 205	6. 139	12, 589	10, 132	7.939	14, 253
	HEMBA1000567								
		41.385	38.483	27. 881	32. 207	15, 544	26.052	15.086	86.601
-	HEMBA1000568	44.686	33.379	126, 524	26. 300	22, 533	17.402	26.970	18.707
5	HEMBA1000569	58. 184	27.187	41.012	21.787	12. 925	36. 191	33.944	23. 225
	HEMBA1000575	155, 833	155.759	434.526	92, 140	79, 143	69.949	59. 928	71. 189
	HEMBA1000588	41.087	26.072	31.610	14. 580	18.024	18.458	23.553	13. 279
	HEMBA1000590	29, 693	17.090	23.618	7. 069	6.633	16.725	20.068	13.042
	HEMBA1000591	106.772	54.874	98. 079					
					34.099	31.776	57.170	48. 488	32.766
10	HEMBA1000592	7, 408	10.031	9. 435	9. 551	8. 209	5.142	7.480	10.319
	HEMBA1000594	18, 401	11.048	22. 547	15. 327	9. 596	12.099	8.751	6.852
	HEMBA1000604	96.047	78.462	146.030	49. 571	36.099	70.815	41.797	47.748
	HEMBA1000607	46.819	15.606	46.037	9. 438	19.149	21.038	17.317	25, 404
	HEMBA1000608	8, 985	3.040	6.705	0.000	7.378	4. 453	0.000	5. 544
	HEMBA1000622	45, 570	55.746	113.666	40. 310	18, 167	19.390	15.895	29.149
	HEMBA1000634	126. 532	49.146	138.073	29.094	95. 787	79.662	60.271	71.657
15	HEMBA1000636		51.270						
		151.899		126. 200	39, 161	51.864	62.611	54.056	39. 415
	HEMBA1000637	33, 241	23.587	39. 380	18.047	16. 265	30.075	28. 225	24. 559
	HEMBA1000655	80.155	70.766	219. 283	58. 901	61, 320	45. 821	40.741	62.639
	HEMBA 1000657	60.961	31.993	41, 401	18.008	30.565	35. 201	35.611	42. 178
	HEMBA 1000662	8, 500	8. 490	11.263	5. 475	2.201	6.140	1.557	2.504
20	HEMBA 1000664	14. 358	5.082	3.637	2.670	3.516	4.913	3.094	3.579
	HEMBA1000671	11, 588	15.473	26.067	17.940	8.865	7.647	10.779	21.196
	HEMBA1000673	73.174	77.410	193. 253	46.051	34, 388	33.975	25.896	31.646 .
	HEMBA1000675	7.666	12.047	22.123	5.764	42.036	15.788	10.254	15. 555
	HEMBA1000678	7.453	12.314	21.083	12.174	14.897	12, 628		
	HEMBA1000682							6.969	6. 584
0.5		118.965	125.696	255.731	86.894	61.443	66. 299	49.060	82.939
25	HEMBA1000686	25.079	17. 463	23.126	12.722	10. 282	13.835	21. 193	18. 154
	HEMBA1000702	206, 683	94. 357	286.585	62.386	79.930	90, 914	98. 397	60.559
	HEMBA1000705	25. 430	25.862	47. 190	13.191	19.599	26. 364	25.013	18.833
	HEMBA1000713	56.893	25. 288	70.751	17.660	24, 138	23, 311	21.805	21.736
	HEMBA1000718	50, 149	43.869	128. 515	28. 289	23.213	18. 458	10.003	17.419
	HEMBA1000719	37.959	17.467	28.513	12.147	12.768	22, 643	14.744	14, 432
30	HEMBA 1000722	15.150	9.762	14.699	6.768	11.726	12.080	5. 907	9. 953
	HEMBA1000726	159.817	111.276	463.937	91.448	109.093	58. 587	46. 517	70.087
	HEMBA1000727	22.867	26.803	28.885	21.475	11.199	14.966	8.634	30. 401
	HEMBA1000732	28.630	11.011	12.790	4.617	3. 548	13. 325	19. 978	13. 472
	HEMBA1000735	24. 568	21.982	21.410	7.431	11.376	41. 026	31.698	
	HEMBA1000743	0.741		1.793					16.801
35			4. 467		1.637	1.227	3.642	4. 563	3.368
30	HEMBA1000745	8.930	7.067	14.546	3.314	10.067	5. 403	9. 225	6.085
	HEMBA1000747	21.442	12.487	25.662	17.081	5. 384	10. 287	9.865	8.267
	HEMBA1000748	22. 924	14.885	35.721	12.634	3, 045	11.508	4.110	11.756
	HEMBA1000749	67. 267	50.826	159.211	43.879	20. 345	29.613	19.447	31.693
	HEMBA1000752	54, 929	35.778	162.005	28. 209	31.540	25. 132	15.650	20.776
	HEMBA1000753	120.889	83.878	155.892	48.092	54, 307	53. 238	38.941	39. 331
40	HEMBA1000757	20. 234	22. 592	52.608	29.935	23.071	24. 503	14.548	43.779
	HEMBA1000760	12, 599	38.665	19.973	15.800	30.188	14, 155	10.570	39. 229
	HEMBA1000769	114.956	74.924	304.424	66.815	39. 365	48, 405	39.918	55. 931
	HEMBA1000773	2, 162	5. 360	11.883	4.445	0.965	3, 158	3.956	2.663
	HEMBA1000774	128. 563	115.732	330.111	84.461	69.618	59, 363	42.656	56. 152
	HEMBA1000780	6.850	7.130	24.175	6.924	6, 903		6.667	9. 576
45	HEMBA1000783						6. 546		
		8. 127	5. 076	13.701	3. 276	8.863	6. 241	5. 435	4. 429
	HEMBA1000791	41.433	51.546	108.542	29.633	42.735	44. 515	43.187	40.856
	HEMBA1000793	108.761	30, 885	54.568	18,670	31.512	54. 669	45. 458	34. 788
	HEMBA1000802	15.062	11, 125	9.052	10.300	11.505	12. 950	15. 354	16. 952
	HEMBA1000813	106.763	52.683	69.701	32.507	44. 369	65.862	59.842	56.799
50	HEMBA1000817	19, 480	7,070	17.915	4.016	15, 239	18.434	11.273	8.079
50	HEMBA1000822	9. 520	10.358	15.760	7.218	8.704	11. 185	6.639	4.662
	HEMBA1000827	96.001	12.420	24.041	8, 305	24.000	6.709	3.488	8.591
	HEMBA1000833	53.675	28, 970	35.897	14, 504	26, 183	29, 036	20, 591	14. 341
	HEMBA1000835	74.696	67.353	83.737	34, 349	42.834	61.145	65.784	52.015
	HEMBA1000843	74.227	54, 197	92.042	37, 825	58. 573	98. 943	87.569	55.077
	HEMBA1000851	23.913	14.070	13.081	6.847	8.634	12.419	19, 200	22. 286
55	HEMBA1000852	56.702	54.074						
				105.085	31, 127	34. 200	31.843	28.843	30. 311
	HEMBA1000867	15, 548	10. 247	11.912	6, 256	1. 227	12. 374	8.518	5. 611

Table 16

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FRIMAN (1008869 19, 566 18, 786 34, 4031 15, 081 6, 290 17, 278 14, 195 17, 765 FRIMAN (1008872 65, 848 46, 546 86, 533 15, 087 40, 538 73, 77 41, 195 37, 195 35, 141 FRIMAN (1008876 38, 976 38, 546 194, 869 47, 595 37, 507 28, 781 17, 455 23, 244 FRIMAN (1008876 38, 976 38, 546 194, 869 47, 595 37, 507 28, 781 17, 455 23, 244 FRIMAN (1008876 38, 976 38, 546 194, 869 47, 595 37, 507 28, 781 17, 455 23, 244 FRIMAN (1008976 47, 197 71, 567 17,							- AAA Y	(A AAA 1	14 206 1	14 718
FERRITIONS 15, 484 16, 546 86, 593 35, 687 40, 506 42, 532 41, 479 36, 141		HEMBA1000869	19.696	18. 785	34.039	15.061	6. 930	13. 298	14. 305	14.712
THE THE		HEMBA1000870	54. 189							
FEBRALDOSS 18, 978 18, 48, 485 42, 355 35, 510 53, 557 36, 311 40, 284 FEBRALDOSS 27, 358 36, 586 10, 377 3, 359 3, 161 53, 557 36, 311 40, 284 11, 587 11, 58	_	HEMBA1000872	46.848	46.546						
	5	HEM8A1000875	35. 460	41.166	32.238	11.297				
		HEMBA1000876	89. 976	56.654	194.869	42.595	57.670			
		HEMBA1000907	22, 959	9,656	10.917	3.599	3.363	5. 327	13.032	10.676
		HEMBA1000908		18, 456	30.665	12, 448	8. 174	19.529	24.789	
						5. 866	7.073	19.938	22.971	11.592
							16.670	25.895	26.769	24.710
	10								23.073	20. 233
HEMBAIDODS25										
HEBBA1000940										
HERBATO00932 125.085 86.817 260.917 52.657 49.118 47.891 15.814 49.81										
15 HEBBAT000945 14, 439 12, 702 14, 590 4, 792 8, 391 11, 856 11, 039 7, 214 HEBBAT009545 17, 850 151, 073 343, 747 107, 319 85, 891 117, 093 82, 928 94, 494 HEBBAT009586 17, 850 151, 073 343, 747 107, 319 85, 891 117, 093 82, 928 94, 494 HEBBAT009586 14, 529 12, 4861 85, 270 8, 173 6, 213 7, 458 7, 214 4, 624 HEBBAT009597 55, 146 19, 281 37, 515 12, 222 19, 562 29, 874 22, 045 23, 135 HEBBAT009571 55, 146 19, 281 37, 515 12, 222 19, 562 29, 874 22, 045 23, 135 HEBBAT009572 51, 003 3, 450 188, 137 28, 972 24, 576 23, 736 11, 32 72, 727 HEBBAT009573 34, 17 19, 192 24, 499 15, 444 40, 09 16, 478 14, 192 14, 153 HEBBAT009575 34, 471 9, 192 24, 499 15, 444 40, 09 16, 478 14, 192 14, 153 HEBBAT009575 35, 417 19, 192 24, 499 15, 444 40, 09 16, 478 14, 192 14, 153 HEBBAT0009581 35, 338 31, 281 33, 672 19, 444 34, 385 38, 280 24, 44, 584 HEBBAT000981 35, 338 31, 281 38, 672 19, 444 34, 385 38, 280 24, 465 31, 286 HEBBAT000983 71, 391 34, 501 58, 683 22, 640 32, 225 32, 384 27, 465 31, 286 HEBBAT000983 72, 707 55, 780 180, 717 34, 576 32, 494 41, 117 32, 433 37, 386 HEBBAT000981 72, 707 55, 780 180, 717 34, 576 32, 494 41, 117 73, 433 37, 386 HEBBAT0000981 72, 707 55, 780 180, 717 34, 576 32, 494 41, 117 73, 43, 33 37, 386 HEBBAT0000981 72, 707 55, 780 180, 717 34, 576 32, 494 41, 117 32, 433 37, 386 HEBBAT0000981 72, 707 55, 780 180, 717 34, 576 32, 494 41, 117 32, 433 37, 386 HEBBAT0000981 73, 77 75, 892 14, 855 85, 565 25, 557 85, 340 53, 718 54, 692 HEBBAT0000981 73, 737 75, 582 14, 855 85, 650 75, 750 85, 640 85, 747 85, 747 85, 748 HEBBAT0000981 73, 737 75, 582 14, 855 85, 650 75, 750 85, 540 85, 543 35,										
HEBBAT000946		77								
HEMBAT000980	15									
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20										
20										
20 HEIBATIOD0372 ST. 100 33,450 188,137 28,972 24,576 23,736 13,731 77,272		HEMBA1000968								
		HEMBA1000971	50. 148							
HEMBA1000975 34.417 19.12 42.499 15.644 4.009 16.478 14.192 14.355 HEMBA1000981 35.318 31.281 38.672 19.544 27.516 38.801 31.347 36.440 HEMBA1000983 371.391 34.501 58.683 22.640 32.625 32.384 27.465 31.286 HEMBA1000986 9.290 20.363 22.497 4.058 6.343 9.035 7.852 3.257 HEMBA1000986 12.114 74.713 236.019 55.652 25.957 85.340 65.711 34.813 38.617 34.815 38.618 38.618 34.815 38.801 37.845 HEMBA1000986 12.114 74.713 236.019 55.652 25.957 85.340 65.711 34.813 37.846 HEMBA1001007 123.690 42.563 59.807 23.525 34.283 47.777 47.496 48.154 HEMBA1001008 37.843 29.269 36.715 11.055 17.115 17.397 17.701 22.055 HEMBA1001009 37.843 29.269 36.715 11.055 17.115 17.397 17.701 22.055 HEMBA10010107 50.408 20.212 48.394 16.20 28.537 31.917 27.876 24.283 HEMBA10010107 73.277 7.582 44.855 6.154 0.598 5.643 3.920 7.188 HEMBA1001020 55.067 55.646 115.814 31.640 25.647 24.596 23.146 27.169 HEMBA1001021 115.724 42.415 59.434 28.288 26.181 54.484 46.173 29.199 HEMBA1001022 23.524 15.235 16.511 8.023 11.818 13.894 3.607 27.169 HEMBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 3.607 29.98 8.047 8.589 HEMBA1001026 71.343 12.515 8.851 6.885 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.288 7.289 7.287	20	HEMBA1000972	51.100	33. 450						
HEBBAT000979		HEMBA1000974	5. 509	10.649						
HEBBA1000981 35, 338 31, 281 38, 672 19,544 34, 383 53, 280 24, 897 29, 059 HEBBA1000985 9, 290 70, 363 22, 497 4, 058 6, 343 9, 035 7, 852 32, 287 HEBBA1000986 128, 714 74, 713 236, 019 55, 662 52, 957 85, 340 6, 3718 54, 892 14, 117 23, 483 37, 845 14, 117 24, 117 23, 483 37, 845 14, 117 24, 117 24, 117 23, 483 37, 845 14, 117 24,		HEMBA1000975	34.417		42, 499	15.644				
THE		HEMBA1000979	90.061	38. 532	99.641		27.516			
PEMBA1000983 71. 391 34. 501 58. 683 22. 640 32. 825 32. 384 27. 465 31. 286 14. 14. 14. 14. 14. 14. 14. 14. 14. 14.			35.338	31.281	38.672	19. 544		38. 280		
PERMATORORS 9,280 20,363 22,497 4,058 6,143 9,015 7,852 3,257 PERMATORORS 128,714 74,713 235,019 56,662 52,957 85,340 63,718 54,882 PERMATORORS 128,714 74,713 235,019 56,662 52,957 85,340 63,718 54,882 PERMATORORS 124,864 47,842 83,746 81,125 25,490 42,631 47,777 47,456 48,154 PERMATORORS 124,864 47,842 83,746 81,125 25,490 52,693 30,688 24,961 PERMATORORS 124,864 47,842 83,746 81,125 25,490 52,693 30,688 24,961 PERMATORORS 124,864 47,842 83,746 81,125 25,490 52,693 30,688 24,961 PERMATORORS 124,864 47,842 83,746 81,125 25,490 26,931 30,688 24,961 PERMATORORS 124,864 47,842 83,746 81,125 25,490 26,931 30,688 24,961 PERMATORORS 124,864 33,356 233,234 60,123 61,977 94,424 47,095 44,625 PERMATORORS 20,212 48,394 16,202 28,537 31,977 27,876 24,283 PERMATORORS 23,067 55,646 115,814 31,640 25,647 24,596 23,146 27,169 PERMATORORS 23,067 55,646 115,814 31,640 25,647 24,596 23,146 27,169 PERMATORORS 23,883 25,835 28,969 18,452 20,270 22,790 25,194 20,783 PERMATORORS 23,832 23,835 28,969 18,452 20,270 22,790 25,194 20,783 PERMATORORS 23,832 21,535 16,511 8,023 11,818 31,834 8,606 8,098 PERMATORORS 23,892 13,860 19,067 12,855 11,445 24,382 15,726 12,323 PERMATORORS 38,892 13,860 19,067 12,855 11,445 24,382 15,726 12,323 PERMATORORS 38,892 13,860 19,067 12,855 11,445 24,382 15,726 12,323 PERMATORORS 33,892 13,860 19,067 12,855 11,445 24,382 15,726 12,323 PERMATORORS 33,892 31,860 19,067 12,855 34,484 33,265 31,860 26,484 PERMATORORS 33,254 26,761 41,713 26,498 24,531 31,498 25,302 23,636 PERMATORORS 47,704 42,537 46,595 31,865 34,686 33,498 55,898 55,574 PERMATORORS 4		HEMBA 1000983	71.391	34. 501	58.683	22.640	32.825	32.384		
HEMBA1000986 128.714 74.713 236.019 56.662 52.957 85.340 63.718 54.892	25		9, 290	20. 363	22.497	4.058	6.343			
HEBBA10010991 72.707 55.730 150.717 34.676 32.494 41.117 23.483 37.846 HEBBA10010007 123.690 42.563 69.807 23.525 34.263 47.777 47.496 48.154 HEBBA1001009 37.843 29.269 36.715 11.055 17.115 17.937 17.701 22.055 HEBBA1001009 37.843 29.269 36.715 11.055 17.115 17.937 17.701 22.055 HEBBA1001014 109.049 83.356 233.234 60.123 61.977.115 17.937 17.701 22.055 HEBBA1001017 50.408 20.212 48.394 16.020 28.537 31.917 27.876 24.283 HEBBA1001019 7.327 7.582 14.865 6.154 10.598 5.643 3.920 7.186 HEBBA1001020 53.067 55.648 115.814 31.640 25.647 24.596 23.146 27.169 HEBBA1001021 115.724 42.415 59.434 28.828 26.181 64.484 64.173 29.614 HEBBA1001022 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEBBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEBBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEBBA1001052 33.892 13.860 19.067 12.855 11.445 24.382 12.663 8.419 7.418 HEBBA1001052 33.892 13.860 19.067 12.855 11.445 24.382 12.663 8.419 7.418 HEBBA1001052 33.892 13.860 19.067 12.855 11.445 24.382 15.726 12.223 HEBBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.223 HEBBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.223 HEBBA1001054 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEBBA1001057 22.4689 23.056 6.665 77.826 26.220 45.125 42.356 36.506 HEBBA1001057 22.4689 23.055 6.665 27.826 26.220 45.125 42.356 36.506 HEBBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEBBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEBBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEBBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEBBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEBBA1001086 77.701 23.951 31.254 22.859 24.531 31.498 25.302 23.636 HEBBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 35.574 HEBBA1001084 5.535 8.834 190.240 540.988 155.496 13.453 93.244 55.99 44.450 17.099 HEBBA1001094 5.535 8.834 190.240 540.	20			74, 713	236.019	56.662	52.957	85. 340	63.718	
			72.707	55. 780	160.717	34.676	32. 494	41. 317		
HEMBA1001008			123,690	42.563	69.807	23. 525	34, 263	47.777	47. 496	48.154
HEMBAIODIOQ9 37.843 29.269 36.715 11.055 17.115 17.937 17.701 22.055 HEMBAIODIO14 109.049 83.356 233.234 60.123 61.977 94.424 47.095 74.625 HEMBAIODIO17 50.408 20.212 48.394 16.020 28.537 31.917 27.376 24.263 HEMBAIODIO19 7.327 7.582 14.865 6.154 10.598 5.643 3.920 7.188 HEMBAIODIO20 53.067 55.646 115.814 31.640 25.647 24.595 23.146 27.169 HEMBAIODIO21 115.774 42.415 59.434 28.828 26.181 64.484 64.173 29.614 HEMBAIODIO22 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEMBAIODIO24 23.524 15.235 16.511 8.023 11.818 33.894 8.606 8.098 HEMBAIODIO24 23.524 15.235 16.511 8.023 11.818 31.894 8.606 8.098 HEMBAIODIO24 23.524 15.235 16.511 8.023 11.818 31.894 8.606 8.098 HEMBAIODIO23 10.374 11.995 9.892 10.750 19.163 9.299 8.047 8.583 HEMBAIODIO53 38.892 33.894 38.006 48.295 68.994 HEMBAIODIO53 38.892 31.850 115.181 387.345 100.376 67.510 61.660 42.295 68.994 HEMBAIODIO52 38.892 31.850 115.181 387.345 100.376 67.510 61.660 42.295 68.994 HEMBAIODIO52 38.897 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEMBAIODIO60 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.612 HEMBAIODIO64 32.254 24.026 33.937 14.007 7.907 33.710 17.387 16.720 HEMBAIODIO77 24.689 23.055 64.486 19.413 16.821 16.858 13.165 12.873 HEMBAIODIO84 32.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBAIODIO85 15.925 24.025 31.937 14.007 7.907 33.710 17.387 16.720 HEMBAIODIO84 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBAIODIO85 159.252 116.909 294.247 77.235 81.384 33.265 31.880 26.484 45.24382 15.726 25.850 14.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBAIODIO84 5.535 8.779 10.059 3.089 4.628 4.521 4.883 4.498 25.302 23.636 HEMBAIODIO93 30.048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEMBAIODIO93 18.6384 190.240 540.908 155.496 134.450 93.524 78.690 116.187 HEMBAIODIO93 18.6384 190.240 540.908 155.496 134.630 93.224 78.890 116.187 HEMBAIODIO104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBAIODIO131 28.384 190.240 540.908 155.496 134.450 93.524 78.690 116.187 HEMBAIODIO122 33.046 52.513				47.842	83.746	18.125	25, 490	52.693	30.668	24. 961
HENBAIODIO14					36.715	11.055	17.115	17.937	17.701	22.055
HEMBA1001017	30				233. 234	60.123	61.977	94. 424		
HEMBA1001019 7. 327 7. 582 14.865 6.154 10.598 5.643 3.920 7.188 HEMBA1001020 53.067 55.646 115.814 31.640 25.647 24.596 23.146 27.169 HEMBA1001021 115.724 42.415 59.434 28.828 26.181 64.484 64.173 29.614 HEMBA1001022 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEMBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBA1001025 21.343 12.515 18.851 6.888 7.288 12.663 8.419 7.418 HEMBA1001051 124.869 115.181 387.345 100.376 67.510 61.660 46.295 68.994 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.327 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.327 HEMBA1001059 98.097 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEMBA1001060 116.837 74.020 161.885 61.750 50.524 52.957 38.575 52.612 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001077 24.689 23.055 54.486 19.413 15.821 16.858 13.165 12.873 HEMBA1001077 24.689 23.055 54.486 19.413 15.821 16.858 13.165 12.873 HEMBA1001077 24.689 27.055 54.486 19.413 15.821 16.858 13.165 12.873 HEMBA1001084 62.698 41.625 171.066 31.438 31.760 24.629 17.487 26.581 HEMBA1001088 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.531 HEMBA1001088 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.531 HEMBA1001089 55.355 8.779 10.059 3.089 4.628 4.521 4.834 4.927 26.531 HEMBA1001089 55.355 8.779 10.059 3.089 4.628 4.521 4.834 4.488 HEMBA1001099 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.488 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001123 18.889 55.587 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001123 18.889 55.587 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001123 18.889 55.587 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001123 18.889 55.587 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367	•				48.394	16.020	28.537	31.917	27.876	24. 283
HEMBA1001020 53.067 55.646 115.814 31.640 25.647 24.596 23.146 27.169 HEMBA1001021 115.724 42.415 59.434 28.828 26.181 64.484 64.173 29.6114 HEMBA1001022 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 14.616 11					14, 865	6.154	10.598	5. 643	3. 920	7. 188
######################################						31.640	25.647	24. 596	23.146	27. 169
HEMBA1001022 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEMBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBA1001025 21.343 12.515 18.851 6.888 7.288 12.663 8.419 7.418 HEMBA1001031 10.374 11.995 9.892 10.750 19.163 9.299 8.047 8.589 HEMBA1001051 124.869 115.181 387.345 100.376 67.510 61.660 46.295 68.994 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001059 93.097 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEMBA1001060 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.612 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001071 25.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBA1001073 23.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBA1001078 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 38.345 59.898 55.574 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 38.34 56.498 29.898 55.574 HEMBA1001085 159.252 116.909 294.247 77.235 38.34 56.498 29.898 55.575 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.468 HEMBA1001095 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA10011091 186.384 190.240 540.908 155.496 134.530 93.324 78.690 116.187 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001133 5						28. 828	26, 181	64.484	64.173	29.614
HEMBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBA1001026 21.343 12.515 18.851 6.888 7.288 12.663 8.419 7.418 HEMBA1001051 124.869 115.181 387.345 10.750 19.163 9.299 8.047 8.589 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001059 98.097 41.525 66.565 27.826 26.220 46.725 42.365 36.506 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001077 24.689 23.055 54.486 19.413 16.821 16.858 13.165 12.873 HEMBA1001080 57.701 23.951 31.254 24.89 24.848 33.265 31.880 26.484 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 55.574 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 55.574 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 55.574 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 55.574 HEMBA1001099 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA10011091 186.384 190.240 540.908 15.46 33.498 44.927 26.310 HEMBA10011093 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA10011093 186.384 190.240 540.908 15.496 134.530 93.224 78.690 116.187 HEMBA10011121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 30.28 42.663 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367							20.270	22.790	25. 194	20. 783
HEMBA1001026 21.343 12.515 18.851 6.888 7.288 12.663 8.419 7.418 HEMBA1001031 10.374 11.995 9.892 10.750 19.163 9.299 8.047 8.589 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001059 98.097 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEMBA1001060 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.612 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001077 24.689 23.055 54.486 19.413 16.821 16.858 13.165 12.873 HEMBA1001078 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.277 77.235 81.384 76.498 59.959 55.574 HEMBA1001086 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBA1001093 30.048 28.810 72.081 20.831 14.610 11.033 15.588 22.531 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.468 HEMBA1001095 18.322 24.021 14.814 7.146 13.778 16.055 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.663 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367	35					8,023	11.818	13.894	8.606	8.098
HEMBAIGOID43 10.374 11.995 9.892 10.750 19.163 9.299 8.047 8.589 HEMBAIGOID51 124.869 115.181 387.345 100.376 67.510 61.660 46.295 68.994 HEMBAIGOID52 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBAIGOID59 98.097 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEMBAIGOID650 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.612 HEMBAIGOID64 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBAIGOID71 25.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBAIGOID78 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBAIGOID78 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBAIGOID88 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBAIGOID88 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBAIGOID88 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBAIGOID89 5.535 8.779 10.059 3.089 4.628 4.521 4.834 44.827 26.310 HEMBAIGOID99 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.468 19.264 13.778 16.055 11.044 10.190 11.059 3.089 4.628 4.521 4.834 4.468 19.264 13.778 16.055 11.044 10.190 11.059 3.089 4.628 4.521 4.834 4.468 11.0100 1000 1000 1000 1000 1000 1000	00						7, 288	12.663	8.419	7.418
HEMBA1001051 124.869 115.181 387.345 100.376 67.510 61.660 45.295 68.994 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001050 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.612 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001071 25.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBA1001073 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.899 55.574 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.899 55.574 HEMBA1001085 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.458 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.468 HEMBA1001095 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001095 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001109 186.384 190.240 540.908 155.496 134.530 93.324 78.6500 116.187 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.268 25.551 30.174 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.4763 34.752 26.702 23.524 11.367							19, 163	9, 299	8.047	8.589
HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001069 98.097 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEMBA1001064 132.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001071 25.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBA1001077 24.689 23.055 64.486 19.413 16.821 16.858 13.165 12.873 HEMBA1001077 24.689 23.055 64.486 19.413 16.821 16.858 13.165 12.873 HEMBA1001078 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 55.574 HEMBA1001093 30.048 28.810 72.081 20.831 14.610 110.33 15.558 22.531 HEMBA1001094 5.535 8.8779 10.059 3.089 4.628 4.521 4.834 4.468 HEMBA1001099 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001109 186.384 190.240 540.98 15.596 134.500 110.33 15.558 22.531 HEMBA1001109 186.384 190.240 540.98 15.597 134.500 193.24 4.500 110.190 186.384 190.240 540.98 15.696 134.530 93.324 78.690 116.187 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.450 21.559 185.527 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.250 9.519 15.621 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.663 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 41.764 34.752 26.702 23.524 11.367							67,510	61,660	46.295	68. 994
HEIBA1001059 98.097 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEIBA1001060 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.612 HEIBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEIBA1001077 24.689 23.055 64.486 19.413 16.821 16.858 13.165 12.873 HEIBA1001078 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEIBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEIBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEIBA1001088 62.698 41.625 171.096 31.438 13.760 24.829 17.487 26.581 HEIBA1001088 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEIBA1001088 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEIBA1001093 30.048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEIBA1001094 5.535 8.779 10.059 30.89 4.628 4.521 4.834 4.468 HEIBA1001094 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEIBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEIBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEIBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEIBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEIBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEIBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEIBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEIBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEIBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEIBA1001121 30.859 55.807 190.789 41.415 39.028 42.663 25.551 30.174										12.323
HEMBA1001060 116.857 74.020 161.885 61.750 50.524 52.957 38.575 52.612 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001071 25.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBA1001078 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.865 12.873 HEMBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.860 26.484 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001088 159.252 116.909 294.247 77.235 81.384 76.489 59.989 55.574 HEMBA1001088 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBA1001083 30.048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.488 HEMBA1001095 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001109 186.384 190.240 540.908 155.496 134.630 93.324 78.690 116.187 HEMBA1001101 186.384 190.240 540.908 155.496 134.630 93.324 78.690 116.187 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.663 25.551 30.174								46,725	42, 356	36.506
HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001071 25.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBA1001078 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.629 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.899 55.574 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.899 55.574 HEMBA1001083 30.048 28.810 77.081 20.831 14.610 11.033 15.558 22.531 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.452 1.688 HEMBA1001095 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001099 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001109 186.384 190.240 540.908 155.496 134.530 93.324 78.690 116.187 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001123 108.859 55.807 190.789 41.4764 34.752 26.702 23.524 11.367	40							52, 957	38.575	52.612
HEIBA1001077 24. 689 23. 055 64. 486 19. 413 16. 821 16. 858 13. 165 12. 873 HEIBA1001077 24. 689 23. 055 64. 486 19. 413 16. 821 16. 858 13. 165 12. 873 HEIBA1001078 33. 254 26. 761 41. 713 26. 498 24. 531 31. 498 25. 302 23. 636 HEIBA1001080 57. 701 23. 951 31. 254 22. 489 24. 848 33. 265 31. 880 26. 484 HEIBA1001084 62. 698 41. 625 171. 096 31. 438 31. 760 24. 829 17. 487 26. 581 HEIBA1001085 159. 252 116. 909 294. 247 77. 235 81. 384 76. 498 59. 989 55. 574 HEIBA1001083 74. 704 42. 537 46. 695 19. 266 25. 146 33. 498 44. 927 26. 310 HEIBA1001093 30. 048 28. 810 72. 081 20. 831 14. 610 11. 033 15. 558 22. 531 HEIBA1001094 5. 535 8. 779 10. 059 3. 089 4. 628 4. 521 4. 834 4. 468 HEIBA1001099 18. 322 24. 021 14. 814 7. 146 13. 778 16. 055 11. 044 10. 190 HEIBA1001109 186. 384 190. 240 540. 908 155. 496 134. 630 93. 324 78. 690 116. 187 HEIBA1001121 32. 820 25. 812 89. 860 19. 710 34. 244 18. 209 9. 519 15. 621 HEIBA1001122 3. 304 6. 213 8. 316 4. 763 19. 120 5. 650 4. 506 4. 506 51. 301 HEIBA1001122 3. 304 6. 213 8. 316 4. 763 19. 120 5. 650 4. 506 3. 059 HEIBA1001123 108. 859 55. 807 190. 789 41. 415 39. 028 42. 663 25. 551 30. 174 HEIBA1001133 50. 744 21. 167 36. 786 14. 764 34. 752 26. 702 23. 524 11. 367	70							13,710	17.387	16.720
HEMBA1001077 24.689 23.055 64.486 19.413 16.821 16.858 13.165 12.873 HEMBA1001078 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEMBA1001084 62.698 41.625 1771.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 55.574 HEMBA1001083 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBA1001093 30.048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.468 HEMBA1001099 18.322 24.021 14.814 7.146 10.10.055 11.044 10.190 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 32.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001114 89.023 252.529 187.547 75.857 35.109 66.259 69.432 341.702 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 39.028 42.663 25.551 30.174 HEMBA1001133 108.859 55.807 190.789 41.415 39.028 42.663 25.551 30.174								9, 530	6.779	24. 242
HEMBAIDDIO78 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBAIDDIO80 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEMBAIDDIO84 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBAIDDIO85 159.252 116.909 294.247 77.235 81.384 76.498 59.595 55.574 HEMBAIDDIO88 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBAIDDIO93 30.048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEMBAIDDIO94 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.488 HEMBAIDDIO994 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.488 HEMBAIDDIO994 5.535 8.779 10.059 3.089 4.628 4.521 4.834 10.190 19.00								16, 858	13.165	12.873
HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.889 55.574 HEMBA1001083 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBA1001083 70.048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.468 HEMBA1001094 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001109 186.384 190.240 540.908 155.496 134.530 93.324 78.690 116.187 HEMBA1001114 89.023 252.529 187.547 75.857 35.109 66.259 69.432 341.702 HEMBA1001121 32.820 25.812 89.860 19.710 34.744 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.663 25.551 30.174										
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HEMBA1001085 159, 252 116, 909 294, 247 77, 235 81, 384 76, 498 59, 989 55, 574 HEMBA1001088 74, 704 42, 537 46, 695 19, 266 25, 146 33, 498 44, 927 26, 310 HEMBA1001093 30, 048 28, 810 72, 081 20, 831 14, 610 11, 033 15, 558 22, 531 HEMBA1001094 5, 535 8, 779 10, 059 3, 089 4, 628 4, 521 4, 834 4, 468 HEMBA1001099 18, 322 24, 021 14, 814 7, 146 13, 778 16, 055 11, 044 10, 190 HEMBA1001104 21, 919 13, 788 35, 048 9, 637 18, 058 24, 450 21, 559 18, 527 HEMBA1001109 186, 384 190, 240 540, 908 155, 496 134, 630 93, 324 78, 690 116, 187 HEMBA1001114 89, 023 252, 529 187, 547 75, 857 35, 109 66, 259 69, 432 341, 702 HEMBA1001127 32, 820 25, 812 89, 860 19, 710 34, 244 18, 209 9, 519 15, 621 HEMBA1001123 108, 859 55, 807 190, 789 41, 475 39, 028 42, 663 25, 551 30, 174 HEMBA1001133 50, 734 21, 167 36, 786 14, 764 34, 752 26, 702 23, 524 11, 367	15									
HEMBA1001088 74,704 42,537 46,695 19,266 25,146 33,498 44,927 26,310	70								59, 989	
HEMBA1001093 30,048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.468 HEMBA1001095 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001109 186.384 190.240 540.908 155.496 134.630 93.324 78.690 116.187 HEMBA10011121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.663 25.551 30.174 HEMBA1001133 50.734 21.167 36.786 14.764 34.752 26.702 23.524 11.367 HEMBA1001133 50.734 21.167 36.786 14.764 34.752 26.702 23.524 11.367										
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HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001109 186.384 190.240 540.908 155.496 134.530 93.324 78.690 116.187 HEMBA1001114 89.023 252.529 187.547 75.857 35.109 66.259 69.432 341.702 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 35.786 14.764 34.752 26.702 23.524 11.367										
50 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527										
HEMBA1001109 186.384 190.240 540.908 155.496 134.630 93.324 78.690 116.187 HEMBA1001114 89.023 252.529 187.547 75.857 35.109 66.259 69.432 341.702 HEMBA1001127 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001123 32.820 33.821 33.821 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 44.145 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367	50									
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HEMBA1001127 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.683 25.551 30.174 55 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367										
HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.683 25.551 30.174 55 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367										
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55 HEMAN 1001 133 50, 744 21.167 36.786 14.764 34.752 26.702 23.524 11.367										
[DEMONICO/131 30.144 21.101 30.700 14.104 04.100 1.000	E E									
HEMBA1001137 38.685 21.659 46.297 21.567 13.174 15.867 11.767 25.508	33									
		HEMBA1001137	38.685	21.659	45.297	21.567	1 13.174	1 15.867	11.767	25. 508

	CONTRACTOR OF THE PARTY OF THE						44 404 1	-0 ACT	42 [61]
	HEMBA1001140	60.453	66.122	169.353	48.837	60.363	44.403	30. 367	43.561
	HEMBA1001144	278.126	195.811		07. 291	166.089	101, 134	105. 337.	142. 120
5	HEMBA1001145	58. 539	241.358	206.084	46.342	39. 316	61.827	91. 170	66.852
3	HEMBA1001158	29.417	28. 121	43.877	13.337	24.176	19.965	18.089	28.622
	HEMBA1001172	74.727	47.695	213.708	37.115	24.460	26.620	19.178	32.709
	HEMBA1001174	6.279	8.617	8.831	7.914	2.574	8.031	3.119	4. 980
	HEMBA1001175	29.561	34.909	43.568	19.819	34. 829	15.588	19.883	17.824
	HEMBA1001182	136.762	64.608	105. 979	44.066	83.417	86,736	126. 297	79.785
			9.703	22.060	9.016	11.018	10. 205	6.347	9, 176
10	HEMBA1001184	16.758					3. 435	9. 089	11. 273
	HEMBA1001192	15.119	10.798	11.626	6.559	5. 736			
	HEMBA 1001197	82.571	114.743	110.687	83.431	56. 396	68.797	99. 959	173. 379
	HEMBA1001208	40. 250	30.964	37. 220	19.514	11.451	24. 172	27.637	12.469
	HEMBA1001213	81.501	37.345	57.618	18.958	24.480	52, 160	51.978	31.326
	HEMBA1001214	36,798	16.011	20.958	17.612	12.418	20.697	19.108	21. 328
15	HEMBA1001221	14, 108	10.456	11.382	7.001	17.058	10.307	7. 980	11.111
13	HEMBA1001225	13.961	14.077	13.384	5.925	5.876	13.456	12.076	5.825
	HEMBA1001226	173. 501	137.685		120.060	113.306	75. 167	63.960	67.304
			48.677	102.518	36.755	64.214	50.002	60.915	35. 178
	HEMBA1001228	115. 971				94. 703	148. 387	156.871	115, 302
	HEMBA1001229	246. 802	111.161,	135.886	43.460				
	HEMBA1001235	43.880	86.102	81.818	36.769	54. 172	65.830	70.065	66. 201
20	HEMBA1001238	67.342	62.561	136.273	36,471	33.652	41.838	26.195	28, 747
	HEMBA1001242	55. 562	43.106	58.593	41.382	47. 200	38, 498	43.114	44. 230
	HEMBA1001247	28.758	22.129	16.518	10.576	8.758	17.031	9.651	13.385
	HEMBA1001253	58. 130	60.415	66.640	18.982	45. 992	54.071	95.073	63.393
	HEMBA1001257	33.557	18.509	24.256	10.657	12.732	31,261	24. 849	9.134
	HEMBA1001261	585. 214	143.415	243.791	98.186	169.988	310.109	234. 388	125.796
25	HEMBA1001262	27. 336	17.339	19.088	5.647	15.678	20.899	11.464	19.889
23	HEMBA1001265	36.604	28.090	152.221	27.730	49.893	34, 423	16.502	26.993
			67.414	170.657	45.898	31.802	39.554	41.287	52.480
	HEMBA1001266	69. 367			34. 126	22. 232	42.207	49.848	39,719
	HEMBA1001269	69. 921	44.649	36.964		8. 604		20.960	13.545
	HEMBA1001272	20. 406	15.416	11.514	7.843		7.893		
	HEMBA1001279	113. 597	76.085	147.371	41.113	50.841	58.248	43, 344	47.548
30	HEMBA1001281	45. 326	37.551	65. 225	44.536	46.787	41.371	32.229	56.625
	HEMBA1001286	370.697	150. 949	236.623	103. 571	123.976	219.461	196.233	117.566
	HEMBA1001289	41.041	24.670	40.151	15.175	30.612	27,627	26.637	19.344
	HEMBA1001291	76.537	40.444	50. 226	18.775	38.423	55. 355	46.692	35.972
	HEMBA1001294	82.258	72.319	157.642	42.143	20.735	29.333	17.711	34.443
	HEMBA1001296	53. 487	17.150	31.045	10.275	15.918	21,120	15.842	13.595
35	HEMBA1001297	13.397	24.306	19.513	11.631	14.701	4.543	9.800	8.121
	HEMBA1001299	122.378	135, 140	326.747	90.817	73.749	56.152	49.803	80.999
	HEMBA1001302	56.839	29.036	56.412	19.108	20,078	34. 481	51.929	37.087
		14. 975	18. 442	43.778	16.797	10. 985	11.442	9,787	19. 264
	HEMBA1001303			244.234	109.949	109. 582	147.334	146.509	115, 543
	HEMBA1001306	262.869	135.864				74. 225	56, 171	58.657
	HEMBA1001308	174.017	96.705	220.049	56.953	61.486			51.006
40	HEMBA1001310	103.029	52.915	67.714	22.895	38. 245	67.233	49. 204	
	HEMBA1001312	98.664	47. 333	61.080	18. 118	33, 555	47.007	41.795	38. 527
	HEMBA1001319	2. 396	8.234	13.960	1.828	5. 485	3.003	5. 682	3.780
	HEMBA1001322	139.794	39.912	105.709	27.700	41,977	70.428	70.602	46.470
	HEMBA1001323	33. 347	16.728	25. 356	11.399	17. 982	11.181	6, 356	12.033
	HEMBA1001326	86.190	37.984	69.933	24. 331	30.078	49.223	46.365	16.347
45	HEMBA1001327	7.232	9.387	23.180	7.314	5. 185	9.563	4. 423	5. 267
	HEMBA1001330	115.768	106.951	275.315	73.389	24.661	70,535	40.088	77.680
	HEMBA1001348	15.770	21.874	26.347	9.575	13.666	23.703	12.647	13.724
	HEMBA1001350		38.749	51.454	16.428	34. 291	56.400	34.055	24.753
		75. 857		56.544	30. 521	46.408	29.604	44.212	30.972
	HEMBA1001351	52. 274	55.313	54.427	17. 559	29. 887	39, 484	52.789	29. 131
· ,	HEMBA1001352	68.321	46.617						
50	HEMBA1001353	39.891	57.492	54.971	31. 425	27.945	45. 587	29.741	66. 188
	HEMBA1001358	45. 659	52. 406	59.774	46.865	40. 225	47.618	32.581	59, 101
	HEMBA1001361	22.908	16.519	28.635	11.897	15.569	13.635	13. 938	16. 914
	HEMBA1001364	18.896	17. 205	23. 355	7. 224	9.469	13.379	76, 125	15. 026
	HEMBA1001375	61.506	22.179	38.795	12.798	25.778	40.077	21.715	22. 300
	HEMBA1001377	140, 430	131.029	307.084	83. 191	100.026	74.475	63. 988	96. 351
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55			26, 206	7 28.704	T 11. 442	17.819	19.160	16.899	7.766
55	HEMBA1001383 HEMBA1001387	23. 974 58. 343	26.206 34.130	28.704 63.677	11. 442		19.160	40. 247	49.239

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	HEMBA1001388	48.601	24.690	39. 877	18. 958	10.634			57.372
	HEMBA1001390	132.003	94.390	254. 352	55.412	64.490	47.169	44. 169	
_	HEMBA1001391	18. 302	9.686	12, 994	6. 299	10.600	8.500	7.116	5. 544
5	HEMBA1001398	91.232	50, 992	142.408	36.081	29.548	29,490	28.704	29. 984
	HEMBA1001405	58.645	22.354	32. 227	15. 864	9. 285	19.993	24. 564	13.964
						19.994	13.316	18.019	16. 592
	HEMBA1001406	36. 434	22,693	105. 808	18.094			18. 826	15.060
	HEMBA1001407	38.781	19.637	24. 599	18. 935	13.107	23,014		
	HEMBA1001411	28. 412	7.180	21.950	8.303	9.708	14.302	8. 598	6.663
	HEMBA1001413	66.736	26.480	35. 635	15, 400	24.013	18.356	24. 304	20.759
10		20.720	7.567	18, 414	12.522	9.722	12.903	18. 283	18.581
	HEMBA1001414					20. 989	32.235	21.694	26.676
	HEMBA1001415	76.802	54.702	159.510	34, 156			25. 847	
	HEMBA1001416	41.784	23, 474	29. 453	12. 230	24.881	24.993		28.651
	HEMBA1001432	74.066	60.077	190.870	40.409	63.619	36.879	66.751	33.675
	HEMBA1001433	132.672	110.163	246.542	77.852	61.676	50.447	37.821	64.403
15	HEMBA1001435	138.669	108, 645	334.104	89. 523	68.855	59.723	58. 393	56. 483
15			8. 504	11, 177	7.985	15.704	7. 291	6.742	6.336
	HEMBA1001442	13. 093				27.976	37. 353	30.266	41.647
	HEMBA1001446	102. 450	63. 255	146, 442	40.085				
	HEMBA1001450	72. 339	35, 494	55, 103	30.799	31.322	42. 457	42.764	41. 349
	HEMBA1001454	148.726	128.060	438.247	88.679	43. 129	54.712	41.131	31.250
	HEMBA1001455	5.879	8, 197	8, 325	5. 561	4. 437	5. 252	4. 300	7.359
20	HEMBA1001459	17.432	15, 927	16, 490	6, 749	2.733	5.888	7.836	10.963
20			52.734	57. 136	38.874	24.764	19.473	23. 241	32.318
	HEMBA1001461	61.531						7.061	25.992
\	HEMBA1001462	10.875	14.911	16. 843	12. 984	13.465	48. 381		
	HEMBA 1001463	137. 907	83.753	340. 496	93.114	51,866	61.784	37.705	68. 960
	HEMBA1001469	85.416	21.757	29. 463	15. 911	84.887	77.440	27. 033	29. 537
	HEMBA1001473	20. 582	31.855	36, 498	8. 307	3, 680	16.703	21.371	19.890
25	HEMBA1001476	135. 720	113, 851	246.800	65. 595	57, 431	63.903	65.229	67.697
20			2.001	4. 505	2.645	1.540	3. 243	1.426	2.876
	HEMBA1001477	5. 228					5. 444	2.171	4. 539
	HEMBA1001478	14. 335	10.180	12,692	5. 468	4.474			
	HEMBA 1001480	88.891	28. 381	49. 689	21.660	14. 126	36. 334	38. 272	30, 563
	HEMBA 1001483	29.872	5. 156	20. 900	4. 647	5. 264	9. 545	13.805	4. 424
	HEMBA1001490	6.867	6. 967	14, 148	7. 289	1.585	5.016	5.792	5. 999
30	HEMBA1001495	431, 282	118,073	203.714	73. 985	176.836	195. 947	194, 164	145. 945
	HEMBA1001497	93.817	60. 807	227.867	55. 576	41.006	34. 182	23.206	45. 223
					75. 008	76. 932	73. 234	61.531	76.899
	HEMBA1001510	174. 254	120, 414	343. 336			10.664	9.567	12.346
	HEMBA1001515	45. 158	26. 337	67. 169	15. 756	15. 962			33. 972
	HEMBA1001517	51,005	47.728	80. 287	34. 595	28. 246	21.020	17.229	
	HEMBA 1001522	7.431	8. 980	7.032	7.566	5.011	8, 466	6.447	4. 824
35	HEMBA1001526	48.774	21.300	32.732	18.831	22.395	22.757	23.530	17. 914
	HEMBA1001533	129. 423	85. 570	262.800	70. 163	46.649	44, 926	26.457	37. 421
	HEMBA1001547	59. 442	26.656	27. 947	8.053	15.558	53, 508	108.861	25. 371
				115. 535	26. 222	30.447	18. 258	21.358	25.853
	HEMBA1001552	41.663	33. 242				54. 513	54.874	41.905
	HEMBA1001553	58, 388	75. 765	66. 228	32.264	36. 396			
	HEMBA1001557	182.516	80. 827	161.852	69.344	80.644	123.765	111.732	70.946
40	HEMBA1001563	39.649	31. 429	85. 246	26.057	12, 157	15. 987	10.065	17.083
	HEMBA1001566	37.835	49.964	108, 284	35.793	23. 255	25. 180	21.368	39. 375
	HEMBA 1001569	75. 584	44. 631	109. 624	35, 487	130, 340	63.130	44.960	55. 257
	HEMBA1001570	198.300	125. 319	444, 153	119, 332	74. 267	79.979	64.732	90.896
			60.654	48.704	22, 469	22.629	67.058	24, 391	34. 300
	HEMBA 1001579	103, 128			131.687	142, 104	91.884	67.267	94.418
45	HEMBA 1001581	153, 698	126, 225	312.570					
40	HEMBA 1001582	3, 551	7.087	15. 302	4.019	8. 190	4. 888	4.671	5. 144
	HEMBA 1001585	27. 271	18. 375	25. 179	14, 108	5.648	14. 993	7.628	12. 297
	HEMBA 1001589	109.877	22.722	49. 216	20. 427	22.904	64.665	57.120	21. 314
	HEMBA1001595	71.600	62.349	46, 938	34. 447	29. 362	34. 516	45. 233	35, 562
			27.004	34, 167	16.004	5.061	21.932	18.414	23. 101
	HEMBA 1001604	41. 253			21.276	22.867	22. 699	14.094	15. 366
50	HEMBA 1001608	35.073	29. 270	41. 525			81.914	42.373	58.652
	HEMBA1001615	556. 575	105. 703	103, 519	47.686	27.311			
	HEMBA 1001620	134, 940	29. 972	79, 824	31.924	62.056	54. 423	64. 359	36.203
	HEMBA1001621	70.036	30.704	63. 807	15.048	19.545	42. 391	33.266	40.516
	HEMBA1001635	39. 932	29. 397	35, 653	16.214	18.765	19.655	22.405	14.095
				35. 798	14. 928	12.865	24. 352	31.819	22.414
	HEMBA 1001636	73.726	18. 596			22, 449	25. 101	30.009	43.819
55	HEMBA 1001640	48. 402	45. 105	79. 588	28. 452				
	HEMBA 1001647	82.402	39. 456	75. 907	35.084	26, 220	48.859	71.158	46. 463
	HEMBA1001651	390, 307	66. 648	181. 929	51,802	112.530	208, 201	178.161	96.640

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				Lac	10 15				
	HEMBA1001655	60.366	18. 983	58. 438	20,404	25. 072	27.162	29. 260	26.673
	HEMBA1001658	6. 754		17. 542	13.420	5.060	4, 800	4. 973	4. 979
			15. 270					22. 916	24. 583
5	HEMBA1001661	87.199	20. 304	32. 793	13.066	8. 394	24. 098		
3	HEMBA1001865	160. 583	20.830	54. 460	12.363	48. 457	86.024	73. 847	21.248
	HENBA1001670	16. 953	38.651	17.002	34, 999	14.855	17.849	22. 906	29. 478
	HEMBA1001672	32, 013	18.885	29.000	10.798	7.753	13.782	17. 314	12.393
	HEMBA1001673	38. 188	67.401	34, 336	38, 037	14, 401	17.612	30. 520	43.461
	HEMBA1001675	25.652	15. 594	33.810	5.190	15, 796	13, 173	20, 020	12, 830
				85. 397	92.681	131.468	50.365	47. 230	68. 405
10	HEMBA1001676	91.000	54.310						
	HEMBA1001678	218.382	128.995	336.408	93.889	115.305	80, 843	48. 879	83. 933
	HEMBA1001680	82.159	51.521	165.818	33.978	35. 449	33.368	38. 495	35. 261
	HEMBA1001681	1.654	0. 785	0.840	2.142	2. 581	2.772	2. 146	2.424
	HEMBA1001684	143.985	84. 151	177.154	72.850	69.097	61.638	30. 820	52.077
	HEMBA1001695	15.068	10.112	14.571	6.860	4. 930	4.572	6. 164	7. 330
	HEMBA1001702	26.509	13.637	8.186	8, 466	4.041	2.043	3.870	3.613
15		67. 279	26. 552	35. 845	13, 982	21.742	28.510	24. 540	19,603
	HEMBA1001709			39, 037	20. 902	21.639	12.713	14.718	33. 127
	HEMBA1001711	20.072	29. 559						
	HEMBA1001712	80.448	25. 222	51.628	19.393	12.482	38.014	39. 474	14. 831
	HEMBA1001714	360.368	55. 902	142. 225	33.748	51.048	144.094	124. 654	59. 543
	HEMBA1001717	78.599	137.380	18. 549	12.298	5. 575	38. 589	10, 120	6.047
20	HEMBA1001718	51.621	52. 280	151. 597	31.305	21, 166	29, 146	14. 075	24.411
	HEMBA1001723	17.072	13.658	8. 525	5.653	8. 811	9.350	11.097	7.268
	HEMBA1001731	35.728	22.781	41.531	15.151	12.421	15. 292	14. 020	16.584
	HEMBA1001734	52, 546	40.599	99. 558	25.099	24.031	28.537	17. 389	32.936
	HEMBA1001736	177. 269	58.328	110,046	33.820	58. 955	108.630	91. 454	62.571
	HEMBA1001741	41. 432	12.649	29. 883	14.886	16, 207	10.446	11. 420	7.286
	HEMBA1001744	5. 531	6.849	12.961	13.191	14, 151	4.519	8. 367	8.623
25				36. 239	12.476	21.118	23.635	15. 410	16, 514
	HEMBA1001745	41.752	17. 786			21. 929	19. 488	11. 306	10.070
	HEMBA1001745	27. 437	14, 874	24, 099	8.668				25. 901
	HEMBA1001761	93. 148	46. 911	179.597	28.212	33. 421	34.026	19. 164	
	HEMBA1001762	55.612	45.069	102, 148	38.307	35.260	33. 316	21. 274	45. 248
	HEMBA1001781	13.298	21.385	26, 693	6.898	17.098	52.601	11.768	23.068
30	HEMBA1001784	89. 965	43.765	70.064	26.575	31.708	50. 347	52. 265	31.618
	HEMBA1001791	182.379	81.719	171.066	44.528	49.350	82.856	58. 215	48. 207
	HEMBA1001794	248. 582	163.789	153.778	73.632	50. 595	152. 279	178.827	132. 329
	HEMBA1001800	23. 432	21.165	27.668	11.281	20.728	24.910	35. 900	22.729
	HEMBA1001803	17, 343	8.333	22.801	6.620	6.043	7.560	6.613	10.079
	HEMBA1001804	109.775	44.797	59.456	29.337	34.849	44. 372	36.696	35.851
35	HEMBA1001808	78. 129	23.567	38.056	15.858	23.507	27.136	14.673	12.332
33	HEMBA1001809	65. 887	31.733	54.127	33.314	26.179	35.618	41.552	46.141
	HEMBA1001811	58, 974	24. 196	37. 583	17.314	16.018	21.582	15.074	19.831
	HEMBA1001815	71. 286	63.775	155, 707	37.153	29.944	35. 297	25. 257	24. 172
				16.797	7.139	5. 598	16.061	22.304	14.646
	HEMBA1001816	38.494	19.017			21. 358	15, 313	14. 917	25. 144
	HEMBA1001819	18, 590	21.371	38. 109	20.938			2. 999	3.099
40	HEMBA1001820	10.884	9.530	8.017	3.507	4. 470	3.473 42.988		47.532
	HEMBA1001822	74. 239	95.719	91.314	62.121	28. 285		38. 222	
	HEMBA1001824	155.543	93, 583	301. 248	95.135	67.478	89.045	64. 562	61.114
	HEM8A1001835	23.616	7.706_	25. 753	5.777	19.660	19.809	12.020	10.462
	HEMBA1001844	149.876	52, 023	230. 213	48.968	42.113	39.652	33.559	40.495
	HEMBA1001847	52.045	19. 220	40.636	20.235	5. 196	35. 109	20. 186	35.814
45	HEMBA1001849	101.048	104, 708	250, 547	53.025	28.022	40.644	33, 371	35. 250
	HEMBA1001850	105. 331	27. 032	39.813	15.808	31.525	42.751	44. 306	18. 213
	HEMBA1001861	3, 104	4. 469	6.763	3.292	4. 454	2.945	0. 995	3.121
	HEMBATDOT862	50. 279	145. 708	102,412	25.750	34.563	40.833	22. 588	71.713
	HEMBA1001864	24.313	31.572	50, 378	32.237	24.991	21.182	21.031	28. 126
			54. 190	146.615	31.714	19. 527	25.041	22.874	21.249
50	HEMBA1001866	57.711					40.562	22.644	40. 891
50	HEMBA1001869	55. 280	99.559	58.454	35.799	45. 195			
	HEMBA 1001871	75.011	44.336	77. 195	41.540	39.300	54. 584	34.598	42.631
	HEMBA1001876	34. 287	31.955	30.568	85.092	19.827	15.356	8. 554	21.861
	HEMBA1001878	17.361	17.619	17.545	15.644	5. 481	11.657	14. 965	18.117
	HEMBA1001879	57.004	22.429	37.128	16.562	20.200	35.414	21. 946	17.114
	HEMBA1001884	68.009	84.640	41.930	38.470	27.460	36.604	25. 345	26.320
<i>5</i> 5	HEMBA1001886	12.711	12.605	37.824	31.827	15.893	14.038	6.697	38.737
	HEMBA1001888	63. 251	46.960	165.623	41.706	21.154	29.117	21. 131	33.090
	(III.MAN 100 1000	1 93.191	1 .0. 500	1,,,,,,,	1, 00				لتتنسب

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	HEMBA1001890	42.902	42.848	42.779	30.112	25. 432	24. 430	22, 605	26.730
			24.720		21.972	17.708	30.703	19.628	23.571
	HEMBA1001896 HEMBA1001899	66. 448	25.553	24. 121	14, 701	12. 301	21.838	17. 455	20.813
5		36. 251			43. 646	55. 587	234, 548	188. 571	526.744
•	HEMBA1001904		10.865	13.738	11. 244	8. 226	15. 367	15. 894	13.300
	HEMBA1001910	40. 309		26. 357	25, 151	11.860	24. 224	22.870	18.238
	HEMBA1001911	35. 962	23. 128		51, 180	45. 903	33, 336	33.019	40.551
	HEMBA1001912	59. 924	66.966	97.679		26. 376	63.459	70.607	52.824
	HEMBA1001913	175. 368	39.664	67. 432	33. 132		17.718	6. 737	8. 522
10	HEM8A1001915	14.756	14.666	30. 224	8. 295	7. 629	8. 265	4. 852	4.772
	HEMBA1001918	5. 018	8.951	27. 591	7.538	11.032	7.417	7.668	2.769
	HEMBA1001921	4, 431	8.444	18.195	11. 252	12. 587	2.935	1. 127	
	HEMBA1001931	3. 948	0.000	4.664	1. 422	3, 480	37.223	29. 835	2.898
	HEMBA1001939	94. 821	24.679	81.706	24, 209	16. 692	18.649	13, 136	19.614
	HEMBA1001940	54. 512	33.931	145. 138	26. 273	27.653		26, 410	
15	HEMBA1001942	38. 572	16.710	32.402	18.718	14. 782	25. 435	92. 097	16. 143
, -	HEMBA 1001944	210,898	71.197	96.883	48. 156	38. 533	82.132		74,740
	HEMBA1001945	31, 531	17.019	14. 533	10. 175	3, 037	17.421	12. 222	11.694
	HEMBA 1001950	7. 103	7.424	9.611	3. 281	4, 091	7.632	5. 310	4.044
	HEMBA1001951	46.024	19. 234	101.026	19. 207	13. 212	23.714	20.006	19.402
	HEMBA 1001958	44. 554	12.806	35. 277	17. 321	13, 181	22.652	28. 735	20.948
20	HEMBA 1001960	20. 513	7.802	16.888	8. 822	2.948	8.826	10.834	12.935
	HEMBA1001962	4. 367	5. 104	4. 205	2.811	3.031	4.870	2. 364	2.994
	HEMBA1001964	35. 944	22, 281	62.751	18.757	6.663	17.775	- 8. 068	8.601
	HEMBA1001967	47.345	29. 504	42.717	13. 526	22. 051	33.555	23.601	37. 521
	HEMBA1001979	35. 138	6, 478	16.732	12.797	5. 919	13.447	10. 355	9.155
	HEMBA1001987	60.083	52.275	190.331	45. 735	24. 898	26.381	17. 514	28.891
25	HEMBA1001991	111.286	79.833	276.566	56. 455	50. 862	50.789	40. 252	54, 919
	HEMBA1002003	66.389	23, 989	51.710	17.039	17, 174	30.547	28. 422	24.474
	HEMBA 1002005	86.885	41,457	150.127	33. 935	15. 339	24. 541	24. 237	27. 345
	HEMBA 1002008	32.101	25. 375	86.511	18. 349	8. 912	7.593	18. 519	14.967
	HEMBA 1002018	66.105	22. 380	36. 174	16. 334	21.482	27.922	34. 098	27.804
	HEMBA 1002022	13, 986	8.018	13. 490	0.000	2.985	5.730	6. 036	1.433
30	HEMBA 1002029	132.547	305.823	115. 974	144. 692	70. 087	74.071	37.046	204. 730
	HEMBA 1002030	17.077	10. 337	14. 524	5. 906	8. 466	5.897	6. 258	6.824
	HEMBA1002035	48. 658	12.959	10.324	14. 325	7.176	14.446	14.084	13.506
	HEMBA1002037	16. 343	34.097	27.567	14. 451	12.568	15.087	13.819	12.140
	HEMBA 1002038	68.477	31.733	91.391	16. 935	8. 370	6.020	17.500	
	HEMBA1002039	15.944	22.707	17.807	13. 914	7.910	3.306	4.716 24.300	22.659
35	HEMBA100204Z	41.657	27.877	32.654	21.111	14, 815	10.217		61.829
	HEMBA1002043	149.364	92.912	208. 642	70.906	53.861	84.089	81. 242 55. 483	30. 137
	HEMBA 1002048	137. 253	29.889	60.279	19.894	21.605	66.594	39. 557	53.676
	HEMBA 1002049	98.417	84.099	271.170	63.157	87. 434 16. 358	16.441	25. 376	27.152
	HEMBA 1002053	33.636	19, 194	25. 821	11.890	17. 449	25. 297	28.606	39.067
	HEMBA 1002055	67, 115	34.916	39.511	37.518 14.136	1.311	8.564	4.481	12.538
40	HEMBA 1002056	13.684	12.039	16.129		10.973	11.179	9.701	8.124
	HEMBA 1002061	11.815	14.960	29. 478 81. 497	10.168	72.416	39.904	45. 653	53.581
	HEMBA 1002080	59.350	80.319 7.502	15. 981	7.301	10.773	13.652	6.835	5. 555
	HEMBA 1002084	11.331	62, 174	111. 196	13.760	19. 083	101.175	43.117	14,011
	HEMBA 1002085 HEMBA 1002092	69.868	33,016	60. 924	24. 219	32.654	72.141	50.433	27.770
	HEMBA 1002092	127. 409	16.695	25. 357	15.741	15.632	18.082	12.882	20, 451
45		34.645	90.733	129. 453	60.276	41.079	89.713	44. 294	67.352
	HEMBA 1002 100	118.301		106.418	34.067	32.565	38. 238	15. 932	74.139
	HEMBA 1002 101	57. 160	69.427 76.058	178.766	45. 801	50. 114	53. 399	40.628	54.459
	HEMBA 1002 102	104.746	25.812	31.300	14. 131	14.867	29.842	22.894	23.960
	HEMBA1002105	35. 380				31.808	52.057		77.437
50	HEMBA1002107	62.621	45.738	65. 486 1335. 986	28. 199	369, 500	391.825	236.013	348.025
50	HEMBA1002113	745.018	396.517		321.385 18.292	19. 991	18. 086	26. 533	25.611
	HEMBA1002119	35, 812	23.546	72.351			47.217	28. 894	33.563
	HEMBA 1002 125	42.106	14.033	45. 440	15,858	20. 474	40.381	49.691	37. 106
	HEMBA1002131	84. 269	29.512	46.944	12.807	29. 311	21.694	20.410	24.305
	HEMBA 1002 133	37.736	19.103	27.034	45. 990	11.161			
66	HEMBA1002139	25.756	10.925	20.941	4. 978	11.839	9.451	7. 795	9.431
55	HEMBA1002141	20.036	14, 349	19.713	9.608	3.638	14.521	10. 225 23. 020	10.190
	HEMBA1002144	86.896	6B, 335	193.756	56,749	45. 612	36.918	23.020	39. 262

	HENBA1002147	135.045	48.848	87. 208	42.412	46.318	67.257	83. 313	45. 988
	HEMBA1002150	347, 113	89.434	182. 502	48.715	86.270	215. 282	234. 394	85. 507
	HEMBA1002151	50,410	19.140	11.868	10.122	7.938	26.996	19.485	14. 196
5	HENBA1002153	32.258	25. 478	35. 746	20. 325	25.638	15. 972	26.019	19.827
	HEMBA1002156	118.226	31.167	44. 382	21, 446	21.743	47. 426	40.620	16.858
				336. 241	90.651	71.047	63.857	41.633	55.419
	HEMBA1002160	166.654	114.853		42. 302	37.035	29. 438	49. 436	41.818
	HEMBA1002161	72.851	68.019	132.156	68. 589	51.141	55. 242	37.823	54. 952
	HEXBA1002162	122,516	52.989	307.464			0.000	30. 979	32. 578
10	HEMBA1002163	49.889	43.602	64. 932	20.426	7. 151	39.440	43. 639	52. 404
	HEMBA1002164	110.221	59.044	71.408	32.652	19. 476		134. 442	
	HEMBA 1002166	312.586	256. 137	758.834	194.638	171.071	159. 271	48. 620	213, 993 13, 548
	HEMBA1002167	139.053	18.430	69.693	11.057	30.091	50. 202		
	HEMBA1002173	137.562	47.918	197.006	36.833	26. 164	29. 478	20. 169	23.078
	HEMBA1002177	100, 895	25. 141	41.676	25.857	17.903	28. 153	22.687	14.081
15	HEMBA 1002178	102.831	19.500	46.717	13.290	32. 323	37.856	44.115	27.390
,-	HEMBA1002179	55.617	56.403	85. 586	45.680	26.918	60. 684	59.110	64. 849
	HEMBA1002185	85. 236	71.958	212.844	43.915	27.049	32.172	22.480	32. 386
	HEMBA1002188	79.413	28. 280	31.826	23.275	21.094	33. 295	36.478	18.236
	HEMBA1002189	56, 349	70.609	148.011	47.092	32.460	30. 101	34, 751	30.532
	HEM8A1002191	149.027	80.765	149.493	49.599	42. 372	50.095	35, 614	44.348
20	HEMBA1002192	15.125	24, 996	24.821	15.373	16. 495	12.778	5.075	13.566
20	HEMBA1002195	57.368	28.635	52. 828	16.254	22. 500	31.843	12. 995	29. 791
	HEMBA1002196 .	14.884	12.040	36.633	16.632	15.443	16.808	12.691	17.451
	HEMBA1002199	24.937	13. 539	27, 878	15.728	17. 426	10. 639	19.664	B. 927
	HEMBA 1002204	9, 525	5. 141	14. 869	6.784	4.619	10. 508	27.818	9.410
	HEMBA1002208	80.832	44, 154	68.317	68.994	37. 453	74.064	81.827	112.820
25	HEMBA1002212	8.709	6, 241	10.946	9.855	2. 602	5.864	5. 366	4.214
20	HEMBA1002215	36. 521	28.098	31.165	19.157	20.170	17. 045	19, 124	21.605
	HEMBA 1002217	50.834	62.759	64.668	59.460	28. 990	37. 379	29. 963	64.813
	HEMBA1002220	27.731	14. 997	21.655	8.451	6. 409	5.663	1.641	6.714
	HEMBA1002226	91.222	113.507	269, 906	85.183	68. 283	59. 461	56.996	78.924
	HEMBA1002227	55.957	91.527	79.169	45. 309	54.892	28.856	14.142	101.597
30	HEMBA1002229	170.518	117, 589	418.739	112.916	121.703	85. 889	63.450	90.668
•	HEMBA1002237	47.252	49. 329	124.721	32.838	24, 807	23. 399	15. 399	26.185
	HEMBA1002239	103.363	107,010	190.830	54.740	72.381	50.451	45. 873	70.581
	HEMBA1002241	70.729	45. 281	81.541	43.824	30. 449	54. 32B	62.401	55.767
	HEMBA1002253	25.559	27.877	35.744	16.605	13.851	18. 938	18. 391	14.286
	HEMBA1002257	6. 344	5.787	15.404	4. 338	1.225	7, 119	4.456	3.711
35	HEM8A1002259	48. 436	19.578	38-228	12.875	21.884	23. 928	18.619	17. 988
00	HEMBA1002262	271.029	219.564	645.284	192.491	147.403	112.552	83.057	137. 280
	HEMBA1002265	56.947	30.786	32.747	24.827	15.078	28.043	29.609	27.237
	HEMBA1002267	108.413	102.522	243.566	58.776	30.097	53.750	24.099	29.752
	HEM8A1002270	51.540	26.396	27.766	20.313	15. 579	28. 348	19.144	16.695
	HEMBA1002286	44.897	17.027	19.776	11.608	10, 900	25. 959	14, 425	10.031
40	HEMBA1002290	46.449	29. 289	34.095	19.879	8.778	26.461	22.368	13. 907 58. 027
,,,	HEMBA1002302	152.883	48, 105	92.158	43.064	48. 204	66.899	80.872	1.737
	HEMBA1002304	8.050	6.814	19.492	7.905	4.038	7.098	5. 307	
	HEMBA1002307	100.402	132.737	29.225	24.612	24.050	42. 355	39.076	37. 573 88. 087
	HEMBA1002316	504.772	93.620	191.534	45.814	134. 386	238.599	265. 167 5. 615	4. 579
	HEMBA1002319	2.868	2, 456	9.670	0.933	4.715	4. 369		3.630
45	HEMBA1002320_	10.783	7. 936	12.646	4.775	10.008	4. 330 7. 952	5. 128 4. 048	5.700
	HEMBA1002321	10.743	9, 992	10.165	4. 549	20. 280	46.772	34.722	18.301
	HEMBA1002328	89.382	28. 578	41.753	17.175			33. 449	21. 452
	HEMBA1002333	63.542	21.208	32.148	11.559	15. 490	29.410	13.954	34. 456
	HEMBA1002337	93.059	61.863	189.067	60.545	43. 745			192.502
	HEMBA1002339	354, 195	154. 586	211.807	141.794	124.733	173. 522 50. 321	284.831 45.600	28. 278
50	HEMBA1002341	116.488	29. 538	63.800	15.812	36.228			
	HEMBA1002348	6.882	4. 859	18.593	4.056	4.011	5. 790		3.748
	HEMBA1002349	5.318	7,600	13.603	5.490	2.590	6.088	1.306	16.760
	HEMBA1002353	14. 497	13.001	12.249	10.426	11.840		17. 141	
	HEMBA1002356	104. 283	29. 278	40.945	24.892	20.681	42. 242		28.190 380.371
	HEMBA1002357	64.855	251.508	219.532	215, 420	68.836	206. 728		
55	HEMBA1002360	87.281	64.882	77.475		55. 108	61.060		56.291
	HEMBA1002363	71.449	51.764	63.278	52.711	43. 280	33, 755	31.248	49. 484

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				140	ne 22			•	
	HEMBA1002365	13.435	10.346	9,534	5.175	9.470	4, 446	10, 802	9. 325
	HEMBA1002370	29. 997	4. 107	11, 054	4.163	3.224	9.009	7,477	2.921
	HEMBA1002374	91.498	18. 475	11. 325	15.862	10.204	18. 275	29. 203	18.856
5	HEMBA1002376	186. 416	75. 425	127.578	52.056	38, 450	99, 590	90, 190	85.994
	HEMBA1002377	81.350	41.908	63, 893	37.221	23.657	110.374	162, 166	50.770
	HEMBA1002380	189. 521	137, 466		137.908	491,500	90, 431	81.778	127.767
	HEMBA1002381	195.037			25.938	88,330	90,756	70, 293	106.965
	HEMBA1002384	35. 247	22.319	42. 496	14.694	19.780	40.126	24. 243	12.399
	HEMBA1002389	44.796	8. 467	36.790	11.793	9.362	18.736	15. 497	20.728
10	HEMBA1002396	101.267	69.467	33.025	16.553	26.429	25. 964	22.294	23.666
	HEMBA1002402	75.818	24. 148	28. 457	8.848	9.913	21, 219	16.569	22.818
	HEM8A1002417	132.807	33, 708	84, 436	22.910	38.826	58, 589	58.836	38. 486
	HEM8A1002419	75, 547	31, 202	41.690	13.558	16.457	27. 281	19.705	13.013
	HEMBA1002420	20.818	20.448	35. 559	17.034	13.878	23.652	14.721	24.637
15	HEMBA1002421	23.903	25. 285	59.023	7.957	14.189	24. 230	61.011	21.849
15	HEMBA1002423	12.762	11.755	25. 941	12.938	14.177	14, 263	12. 495	7.512
	HEMBA1002424	111. 995	32. 293	45. 657	24. 424	25.667	42.797	41.513	31.249
	HEMBA1002426	60.617	23. 489	45. 906	20.305	25.173	30, 860	37.738	21.223
	HEMBA1002430	24. 143	3. 1284	4. 900	1.517	4.594	3. 316	8. 552	3.069
	HEMBA1002439	59. 808	37, 476	93. 025	16.789	23.324	47.857	33.099	27.888
20	HEMBA1002441	77.869	99. 262	110.341	38.723	34.562	65. 109	85.421 22.845	66. 581 22. 015
	HEMBA1002454	58. 292	15. 281	38. 384	7. 520	19.044 82.184	25.972 61.800	26.094	59.039
	HEMBA1002458	57, 329	46. 103	25. 085	30, 906 12, 160	23.009	18,683	14.678	14, 249
	HEMBA1002460	12.814	9. 205 38. 135	55. 208	10. 919	24. 257	49.697	43.851	32. 387
	HEMBA1002462 HEMBA1002465	98. 420 11. 819	15. 260	28. 272	11. 939	11.225	10. 938	13.593	20.635
	HEMBA1002469	129. 538	61.348	120. 187	39. 999	39.213	76, 320	69.012	86.309
25	HEMBA1002475	3.180	5, 116	4. 323	2.230	1.467	4. 495	10.058	15.691
	HEMBA1002477	93.696	64.730	238, 114	55. 207	43, 349	42.487	29. 532	52.786
	HEMBA1002480	210.023	58.823	84. 566	37.478	45.060	106.554	97.791	70.487
	HEMBA1002481	104. 499	76.474	222. 903	71.502	68.097	67.421	42. 334	82.875
	HEMBA1002486	81.465	42.269	169. 291	49. 953	40.852	39. 475	29. 153	26. 233
30	HEMBA1002490	66. 695	11. 331	31.314	14.602	25.852	35.945	35, 954	15. 278
	HEMBA1002495	59.387	12, 315	25. 235	7, 937	4, 091	17.402	14. 269	10.773
	HEMBA1002498	56.425	23.969	67. 108	11.632	15.655	24.420	8. 272 39. 506	12.219 24.619
	HEMBA1002501	40.955	16. 994	22.074	13. 575	16.498 33.778	31.214	32.219	26. 800
	HEMBA1002503	81.763	65.044	154.595 279.391	39. 638 90. 092	120. 246	70.516	52.190	53.323
	HEMBA1002504	155. 357	95. 219 88. 234	259. 961	107. 085	79.039	59. 181	59. 924	61. 423
35	HEMBA1002508 HEMBA1002513	50, 560	22. 902	30. 431	26. 184	20.783	30, 500	32.903	22.864
	HEMBA1002515	60.938	23.064	25. 098	16. 172	5.716	20.264	20.643	13.727
	HEMBA1002524	94. 350	36.789	56.675	25. 998	28.978	49.840	57. 148	25. 205
	HEMBA1002538	116,609	19.632	26.764	12.798	20, 203	15.422	17. 588	15.759
	HEMBA1002542	81.641	81.952	188.888	54. 986	41.864	32.890	30.719	38. 321
40	HEMBA1002544	52.394	49.175	98. 415	47.569	28.375	28, 766	20, 948	21.614
70	HEMBA 1002546	76.538	62.763	156.051	47.625	74.374	45.975	34.756	46.753
	HEMBA1002547	11.448	4.516	10.647	4. 733	12, 220	11.801	9, 959 94, 586	7. 127
	HEMBA 1002550	67.373	39. 322	48.468	15.671	16.497	121.814	16.656	25. 401 18. 665
	HEMBA1002551	94. 391	14. 109	27.085	11. 976 49. 448	8. 787 44. 756	67.408	63.216	57.684
	HEMBA1002552	204. 583	77.430	205. 444 6. 743	7. 020	5.608	14.795	10.111	7.416
45	HEMBA1002555	25. 583	16. 987 77. 405	245, 703	59.079	41.247	33.253	41.617	41.270
	HENBA1002558	92.744 53.810	51.725	155. 895	34. 956	27.689	17.264	10.138	27.124
	HENBA1002562	15, 261	10.822	15. 435	8. 259	18.723	12,036	9, 056	10. 429
	HEMBA1002568	24, 946	17.442	35. 354	17.552	10.576	15. 262	16, 158	22.328
	HEMBA1002569	112, 340	34, 133	118, 192		57, 431	54.936	26.164	27. 309
50	HEMBA1002570	43. 528	50.809	52. 195			28.874	9.812	50. 494
~	HEMBA 1002574	106, 101	25.148	46.793			57.278	42. 795	31.310
	HEMBA 1002583	36.042	17.582	15, 178		13.418	20.158	16.837	16.418
	HENBA 1002587	61.527	32. 123	45.811			32.461	40. 250	39. 915
	HEMBA1002590	151.583	106.074	287.276			58. 221	38, 542	53.855
	HEMBA1002592	97.854	85. 949	220. 496			53.653	35.724	
55	HEMBA 1002595	146.016	25. 688	60. 427			86.770	79.174	
	HEMBA 1002609	97.442	41. 926	56,054	29. 427	35.650	35.839	44.688	47.074

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HEMBAIRODEST 12, 127			*****			40 544 1	10 000 1	21 625 1	10 614 1	49, 136
FERRALITIONS 14, 552 15, 345 16, 082 16, 272 7, 391 18, 565 5, 522 12, 283 18, 1861 19, 1871 19, 125 19, 1871 19, 1872 19, 1872 19, 1874 19, 1		HEMBA1002617	26.792	86.617	59. 446	73.277	12.909	21.055	16.612	
FERBAIDOVERS 14, 552 25, 815 18, 082 8, 927 7, 391 8, 869 5, 523 12, 283 FERBAIDOVERS 19, 044 21, 509 31, 449 9, 986 16, 559 10, 257 7, 527 11, 624 FERBAIDOVERS 27, 199 16, 370 27, 140 3, 985 16, 559 10, 257 7, 527 11, 624 FERBAIDOVERS 32, 199 86, 370 27, 140 15, 884 3, 171 46, 444 15, 444 15, 585 14, 644 15, 644		HEMBA1002619	101, 131	25, 998	30.959	15.921	21.913	40.814	35.003	
Second property 254, 5515 27, 287 73, 558 48, 038 71, 672 113, 228 101, 788 53, 514 FEBRI 10071279 32, 198 16, 370 37, 306 15, 884 5, 559 10, 275 75, 527 11, 624 FEBRI 10071279 32, 198 16, 370 37, 306 15, 884 5, 272 15, 410 42, 984 19, 880 FEBRI 10071279 32, 198 16, 370 37, 306 15, 884 5, 272 15, 410 42, 984 19, 880 FEBRI 10071279 32, 198 16, 370 37, 306 15, 884 5, 272 15, 410 42, 984 19, 880 FEBRI 10071279 32, 198 16, 370 37, 306 15, 884 5, 272 15, 410 42, 984 19, 880 FEBRI 10071279 32, 198 16, 370 37, 313 16, 598 17, 48, 643 56, 847 41, 355 59, 867 FEBRI 10071279 37, 307 37, 307 37, 307 37, 307 37, 216 28, 227 37, 227 FEBRI 10071279 37, 307 3						8 927	7, 391	8 869	5, 823	12, 283
HEBBA10075129 37.198 18.770 27.306 38.58 16.559 10.257 7.527 11.624 11.62	_									
THE SELF HORSE 15 15 15 15 15 15 15 1	5	HEMBA1002624								
		HEMBA1002528	13.044	21.509	23.649	9.956	16.559	10.257	7.527	
REMINITORISTS 78, 578 48, 044 90, 986 81, 904 27, 840 28, 811 37, 912 40, 948					29 306	15 884	5, 722	15,410	42, 964	19.680
		HEMBA 1002632								
		HEMBA 1002645	95, 909	89.897	220.184	68, 171	48.643			
					33 313	16.958	11,617	23, 904	29, 214	16, 599
HEBBA1002785 331, 320 62, 916 259, 854 57, 860 53, 172 46, 511 45, 193 47, 291 HEBBA1002786 38, 436 48, 41, 70 35, 198 22, 499 26, 290 22, 41 23, 727 HEBBA1002786 38, 174 70, 511 39, 391 77, 036 15, 852 20, 842 19, 202 13, 470 14, 170 15, 185 23, 185 24, 1	10							37 216		
HEBBA1002786 38.495 68.014 154.170 35.196 22.499 22.210 22.314 23.727 HEBBA1002787 75.5384 156.224 164.588 29.523 520.013 30.234 25.612 33.769 HEBBA1002787 151.581 89.986 247.534 84.727 84.017 440.179 318.619 22.512 HEBBA1002787 161.581 89.986 247.534 84.727 84.176 46.941 61.944 77.055 HEBBA1002787 161.581 89.986 247.534 84.727 84.176 46.941 61.944 77.055 HEBBA1002788 6.756 3.564 55.37 37.679 18.172 29.420 38.288 44.113 HEBBA1002789 49.619 17.555 29.241 14.788 17.93 4.608 3.600 2.944 HEBBA1002706 45.533 30.340 35.619 18.172 29.420 38.288 44.113 HEBBA1002707 45.538 96.718 77.793 54.473 55.688 113.880 87.727 59.377 HEBBA1002707 45.538 96.718 77.793 54.473 55.688 113.880 87.727 59.377 HEBBA1002708 45.533 30.340 35.619 18.469 19.18 26.777 79.277 79.272 59.378 HEBBA1002706 45.533 30.340 35.619 18.469 19.18 26.777 79.277 79.272 HEBBA1002707 52.528 59.111 10.505 41.591 42.597 42.591 77.808 55.80 HEBBA1002715 149.045 59.588 87.643 47.473 41.264 59.579 17.808 55.80 HEBBA1002716 25.328 19.063 41.749 25.345 16.735 28.367 26.822 21.779 HEBBA1002778 17.388 88.550 27.971 88.69 19.18 25.577 19.644 77.572 HEBBA1002778 17.588 88.550 27.971 28.681 28.385 49.987 50.300 43.793 HEBBA1002774 10.730 11.726 12.788 79.19 13.948 89.51 17.955 89.487 19.487 89.51 19.644 27.647 75.72 HEBBA1002774 75.748 28.513 38.699 17.609 19.194 89.51 19.594 19.644 27.547 45.992 HEBBA1002774 75.748 27.585 27.205 28.657 18.700 28.233 31.497 33.333 18.347 HEBBA1002774 75.748 27.585 27.205 28.657 18.224 37.555 37.707 27.737 37.747 37.77 37.77 37.77 37.77 37.77 37.77 37.77 37.77 37.77 37.77 37.77 37.77 37	10	HEMBA1002652						-10 F11		
HEBBA1002786	•	HEMBA1002659	133. 320	62.916	259.854	57.860	53.172	46.511		
HEBBA1002FSF 75.34 16.24 19.202 13.470				68 014	154, 170 I	35, 196	22.499	26. 290	22.314	23. 727
HEMBATO02787 155. 384 156. 224 164. 586 29. 523 320. 013 30. 214 25. 612 83. 765 HEMBATO02787 161. 681 88. 386 247. 534 84. 722 54. 175 46. 941 61. 944 77. 085 HEMBATO02787 161. 681 88. 386 247. 534 84. 722 54. 175 46. 941 61. 944 77. 085 HEMBATO02787 161. 681 88. 386 247. 534 84. 722 54. 175 46. 941 61. 944 77. 085 HEMBATO02787 55. 416 61. 838 68. 537 37. 679 81. 29. 340 32. 238 44. 113 HEMBATO02788 6. 756 3. 164 5. 387 37. 679 81. 29. 340 38. 238 44. 113 HEMBATO02708 48. 533 96. 718 97. 739 54. 473 56. 688 81. 380 87. 727 59. 878 HEMBATO02706 48. 533 30. 340 35. 879 18. 469 19. 118 26. 777 79. 277 29. 272 REMBATO02706 48. 533 30. 340 35. 879 18. 469 19. 118 26. 777 79. 277 29. 272 HEMBATO02706 48. 533 30. 340 35. 879 18. 469 19. 118 26. 777 79. 277 29. 272 HEMBATO02715 149. 045 59. 358 87. 643 47. 473 41. 264 95. 279 17. 808 65. 801 HEMBATO02716 23. 248 88. 550 27. 71 70. 77 15. 832 23. 557 19. 644 27. 647 7. 572 HEMBATO02718 17. 868 88. 550 37. 091 81. 269 95. 279 17. 808 65. 801 HEMBATO02718 17. 868 88. 550 27. 97. 18. 628 36. 688 49. 987 50. 360 43. 208 HEMBATO02718 17. 868 88. 550 67. 877 28. 628 36. 688 49. 987 50. 360 43. 208 HEMBATO02718 17. 868 27. 862 67. 877 28. 628 36. 688 49. 987 50. 360 43. 208 HEMBATO02718 10. 330 11. 726 12. 788 79. 19 13. 194 8. 502 8. 297 10. 599 HEMBATO02718 10. 330 17. 776 12. 788 79. 19 13. 94 8. 502 8. 297 10. 599 HEMBATO02718 10. 300 17. 776 12. 788 79. 19 13. 194 8. 502 8. 297 10. 599 HEMBATO02718 10. 300 37. 554 59. 500 38. 215 37. 19. 19. 19. 19. 19. 19. 19. 19. 19. 19								20 842	19, 202	13 470
		HEMBA1002667								
HEBRATODZ\$F9		HEMBA1002673	71, 650	40.718	73.822	33.403	39.914	40.129	38, 619	22.532
### FEBRAT002879	15					84 722	54, 176	46, 941	61.944	77.085
HEBRIDOZESS 3,756 5,364 5,387 3,816 1,793 4,608 3,600 2,944 HEBRIDOZESS 49,819 17,555 29,241 14,788 12,483 31,752 34,100 14,772 HEBRIDOZTOS 185,328 96,718 97,733 54,473 50,688 113,980 87,727 59,878 HEBRIDOZTOS 49,533 30,340 35,679 18,489 19,118 26,777 29,277 29,277 29,277 HEBRIDOZTIS 25,878 59,111 110,506 41,591 43,597 39,604 217,28,224 HEBRIDOZTIS 149,045 59,858 87,633 47,473 41,264 95,279 127,808 65,580 HEBRIDOZTIS 149,045 59,858 87,633 47,473 41,264 95,279 127,808 65,580 HEBRIDOZTIS 12,045 59,858 87,633 47,473 41,264 95,279 127,808 65,580 HEBRIDOZTIS 17,984 88,850 231,019 81,759 43,779 55,830 62,121 57,003 HEBRIDOZTIS 117,784 88,852 637,019 81,290 43,779 55,830 62,121 57,003 HEBRIDOZTIS 117,784 88,852 637,817 28,628 86,666 49,877 50,300 43,208 HEBRIDOZTIS 17,787 26,481 34,604 21,128 21,756 49,877 55,830 43,208 HEBRIDOZTIS 40,653 43,306 95,205 18,200 10,037 22,527 29,311 30,774 HEBRIDOZTIS 40,653 45,306 95,205 18,200 10,037 22,527 29,311 30,774 HEBRIDOZTIS 40,653 45,306 95,205 18,200 10,037 22,527 29,311 30,774 HEBRIDOZTIS 40,653 45,306 95,205 18,200 10,037 22,527 29,311 30,774 HEBRIDOZTIS 40,653 45,306 95,205 18,200 10,037 22,527 29,311 30,774 HEBRIDOZTIS 40,653 45,306 95,205 18,200 10,037 22,527 29,311 30,774 HEBRIDOZTIS 40,653 45,306 95,205 18,200 10,037 22,527 29,311 30,774 HEBRIDOZTIS 40,593										
HEIBATIODZ755										
HEMBA1002703 185. 328 56. 718 97. 793 54. 473 50. 688 113. 980 87. 727 59. 878 HEMBA1002705 49. 533 30. 340 35. 679 18. 468 19. 118 26. 777 29. 224 HEMBA1002715 149.045 59. 858 87. 643 47. 473 41. 264 95. 279 127. 808 65. 580 HEMBA1002716 149.045 59. 858 87. 643 47. 473 41. 264 95. 279 127. 808 65. 580 HEMBA1002718 26. 328 19. 663 41. 749 26. 345 16. 715 28. 367 26. 822 21. 779 HEMBA1002718 26. 328 19. 663 41. 749 26. 345 16. 715 28. 367 26. 822 21. 779 HEMBA1002728 117. 848 88. 550 293.019 81. 290 43. 679 65. 830 44. 221 57. 603 HEMBA1002734 77. 679 26. 481 34. 604 21. 128 21. 755 41. 413 60. 557 45. 982 HEMBA1002742 77. 679 26. 481 34. 604 21. 128 21. 755 41. 413 60. 557 45. 982 HEMBA1002746 60. 876 22. 803 35. 400 15. 330 15. 630 30. 605 31. 889 32. 759 HEMBA1002746 60. 876 22. 803 35. 400 15. 330 15. 630 30. 605 31. 889 32. 759 HEMBA1002755 94. 758 62. 505 270. 964 63. 144 37. 572 24. 593 28. 497 39. 377 HEMBA1002755 94. 758 62. 505 270. 964 63. 144 37. 572 24. 593 28. 497 39. 377 HEMBA1002767 60. 873 62. 505 270. 964 63. 144 37. 572 24. 593 28. 497 39. 377 HEMBA1002767 74. 568 26. 505 270. 964 63. 144 37. 572 24. 593 28. 497 39. 377 HEMBA1002767 103. 615 37. 655 72. 072 41. 794 31. 219 54. 881 59. 342 41. 525 HEMBA1002767 103. 615 37. 655 72. 072 41. 794 31. 219 54. 881 59. 342 41. 525 HEMBA1002777 23. 615 23. 565 27. 877 27. 772 27. 773 27. 770 27. 264 39. 30. 30. 30. 30. 30. 30. 30. 30. 30. 30		HEMBA1002688	6.756	3.364	5. 38 <u>/</u> _					
HEIBAT10027705		HEMRA 1002696	49 639	17.555	29, 241	14.788	12.463	31.752	34, 100	14.772
HEMBA1002775								113 980	87 727	59.878
HEBBA1002715	00	HEMBA1002706								
HEBBA10027716	20	HEMBA1002712	52, 878	59. 111	110.506	41.591	43.597	39.604	30.872	26.457
HEBBA10027716 23.142 6.155 17.077 15.183 23.557 19.064 27.647 7.572 HEBBA10027718 75.288 19.053 41.749 26.345 16.735 28.367 25.822 21.779 HEBBA10027730 131.726 26.862 67.877 28.623 36.686 49.987 25.380 43.208 HEBBA10027742 10.730 11.276 26.862 67.877 28.623 36.686 49.987 25.0380 43.208 HEBBA1002744 10.730 11.276 12.768 7.910 1.394 8.502 8.297 10.909 HEBBA1002745 60.876 22.803 35.400 15.830 15.630 30.605 31.889 32.759 HEBBA1002755 40.653 45.306 95.205 18.200 10.377 22.527 29.31 30.774 HEBBA1002755 94.758 62.505 220.964 63.444 37.572 44.593 28.497 39.737 HEBBA1002755 43.094 88.931 172.920 71.623 88.921 195.471 197.995 118.224 HEBBA1002767 50.682 25.272 33.782 40.358 195.471 197.995 118.224 HEBBA1002768 100.803 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEBBA10027769 03.210 30.236 34.098 17.769 31.893 31.497 33.193 38.347 HEBBA10027770 97.457 29.259 75.005 22.719 22.637 33.693 38.371 HEBBA10027780 72.338 50.411 181.355 42.070 19.957 31.350 34.931 35.440 HEBBA10027780 72.338 50.411 181.355 42.070 19.957 31.370 27.642 39.672 HEBBA10027780 72.338 50.411 181.355 42.070 19.957 31.370 27.642 39.672 HEBBA10027780 72.338 50.411 181.355 42.070 19.957 31.370 27.642 39.672 HEBBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672 HEBBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672 HEBBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672 HEBBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672 HEBBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672 HEBBA1002786 72.538 50.411 181.356 42.070 19.957 31.370 27					87 643	47.473	41.264	95, 279	127, 808	65, 580
HEIBA 1002718										
		HEMBA1002718	26.328	19.063						
HEBBA1002730 131.726 26.862 67.877 28.628 36.686 49.987 50.380 43.208 HEBBA1002744 77.679 26.481 34.604 21.128 21.756 41.413 60.657 45.992 HEBBA1002746 60.876 72.803 35.400 15.830 15.630 30.605 31.889 22.759 HEBBA1002746 60.876 72.803 35.400 15.830 15.630 30.605 31.889 22.759 HEBBA1002746 60.876 72.803 35.400 15.830 15.630 30.605 31.889 22.759 HEBBA1002748 76.748 25.130 38.669 17.760 32.833 43.493 51.440 49.691 HEBBA1002750 40.653 45.306 95.205 18.200 10.037 22.527 29.331 30.774 HEBBA1002755 94.758 62.505 270.964 61.414 37.572 44.593 28.497 39.737 HEBBA1002767 43.935 31.17 8.450 37.92 2.291 8.714 10.261 5.285 HEBBA1002763 430.941 88.931 172.920 71.623 88.921 195.471 197.995 118.224 HEBBA1002767 65.682 25.272 35.782 14.035 19.83 31.497 33.393 83.347 HEBBA1002769 103.210 30.236 54.098 77.099 19.753 35.636 41.922 25.940 HEBBA1002769 103.210 30.236 54.098 77.099 19.753 35.636 41.922 25.940 HEBBA10027767 40.615 37.655 72.072 41.794 31.219 54.881 59.342 43.652 HEBBA1002779 97.457 29.259 75.705 22.719 22.643 33.699 38.357 77.804 HEBBA1002779 97.457 29.259 75.705 22.719 22.643 33.699 38.357 77.804 HEBBA1002798 9.194 21.334 22.488 20.353 79.745 39.803 38.357 77.804 HEBBA1002780 77.338 59.411 181.356 24.070 19.575 31.370 27.642 30.672 HEBBA1002781 42.583 45.313 55.084 37.799 84.893 81.695 152.339 171.186 177.409 HEBBA1002807 139.924 07.778 533.137 90.533 79.745 59.859 54.005 59.584 HEBBA1002807 139.924 07.778 533.137 90.533 79.745 59.859 54.005 59.598 HEBBA1002808 77.126 30.401 44.872 22.577 28.63 50.244 43.98 57.302 35.668 HEBBA1002808 77.126 30.401 44.872 22.577 28.63 5		HEMBA1002728	117, 984	88, 950	293.019	81.290	43.679	65.830	46. 321	57.003
HEBBA10027734 77.679 26.481 34.504 21.128 21.755 41.413 50.057 45.992 HEBBA1002742 10.730 11.276 12.768 7.910 1.394 8.502 8.297 10.909 HEBBA1002748 76.748 25.130 38.669 17.760 32.833 43.493 53.440 49.691 HEBBA1002745 76.748 25.130 38.669 17.760 32.833 43.493 53.440 49.691 HEBBA1002755 94.758 67.505 270.964 63.414 37.572 44.593 28.497 39.737 HEBBA1002755 94.758 67.505 270.964 63.414 37.572 44.593 28.497 39.737 HEBBA1002767 40.941 88.931 17.290 71.623 88.921 195.471 19.541 19.795 118.244 HEBBA1002767 65.682 25.272 35.782 14.035 19.183 31.497 33.393 18.347 HEBBA1002767 65.682 25.272 35.782 14.035 19.183 31.497 33.393 18.347 HEBBA1002767 60.083 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEBBA1002777 103.210 30.236 54.098 17.099 19.53 35.636 41.922 28.900 HEBBA1002777 20.350 15.268 28.054 21.735 10.754 12.030 14.991 11.776 HEBBA1002778 77.336 55.72 77.2 41.794 31.219 54.881 59.342 43.652 HEBBA1002779 20.350 15.268 28.054 27.719 27.643 33.689 38.357 27.804 HEBBA1002779 87.377 61.291 152.514 38.033 29.564 28.032 20.352 34.761 HEBBA1002799 87.377 61.291 152.514 38.033 29.56 28.032 20.352 34.761 HEBBA1002799 89.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEBBA1002799 89.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEBBA1002798 77.751 59.583 33.317 90.533 79.745 59.869 54.302 52.952 HEBBA1002798 79.947 27.383 45.313 55.086 35.416 29.480 60.935 44.046 51.794 HEBBA1002810 42.583 45.313 55.086 35.416 29.480 60.935 44.046 51.794 HEBBA1002826 40.776 6.495 6.825 5.349 31.90 4.400 3.305 57.305 36.688 HEBBA1002863 77.126 30.407 44.248 40.339 77.845 23.324							36,686	49, 987	50, 380	43, 208
HEBBA1002748	25									
FEMBA 1002748 60.876 22.803 35.400 15.830 15.630 30.605 31.889 32.759 FEMBA 1002748 76.748 26.130 38.669 17.760 32.833 43.493 53.440 49.691 FEMBA 1002755 40.663 45.306 95.205 18.200 10.37 22.527 29.331 30.774 FEMBA 1002755 94.758 62.505 220.964 63.414 37.572 44.593 28.497 39.737 FEMBA 1002759 13.935 3.117 8.450 3.792 2.291 8.714 10.261 5.285 FEMBA 1002767 65.682 25.272 35.782 14.035 19.183 31.497 33.393 18.347 FEMBA 1002768 100.803 57.554 59.457 35.570 28.006 37.70 40.930 38.215 FEMBA 1002769 103.210 30.236 54.098 17.099 19.753 35.636 41.922 26.940 FEMBA 1002770 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776 FEMBA 1002777 130.615 37.655 72.072 41.794 31.219 54.881 59.342 43.652 FEMBA 1002779 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 FEMBA 1002780 77.338 50.411 181.356 42.070 19.957 11.370 27.642 39.672 FEMBA 1002780 87.371 61.291 152.514 38.033 29.516 28.032 20.352 34.761 FEMBA 1002788 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 FEMBA 1002780 42.533 45.313 55.088 35.416 29.480 60.935 44.046 51.794 FEMBA 1002818 371.516 100.826 187.799 84.893 81.595 152.339 711.86 117.405 FEMBA 10028286 40.776 6.495 16.825 5.499 3.590 5.574 33.389 50.974 49.045 FEMBA 1002826 60.735 32.524 30.030 9.693 9.527 27.595 19.397 18.101 FEMBA 1002866 77.126 30.401 44.288 40.839 7.555 7.855 59.859 54.302 52.958 FEMBA 1002866 77.126 30.401 44.872 22.577 28.633 30.995 15.800 30.566 38.689 FEMBA 1002866 77.126 30.401 44.872 22.577 28.633 30.995 39.971 49.045 FEMBA 1002866 77.126 30.401 44.872 22.577 28.633 30.995 57.302 36.668 FEMBA 1002886 9.474 14.188 23.688	20									
HEMBA10027748		HEMBA1002742	10.730	11.276	12.768	7.910	1. 394			
HEMBA1002748 76.748 25.130 38.669 17.760 32.833 43.493 53.440 49.691 HEMBA1002750 40.663 45.306 95.205 18.200 10.037 22.527 29.331 30.774 HEMBA1002759 34.758 62.505 270.964 63.414 37.572 44.593 28.497 39.737 HEMBA1002759 13.935 3.117 8.450 3.792 2.291 8.714 10.261 5.285 HEMBA1002763 430.941 88.931 172.920 71.623 88.921 195.471 197.995 118.224 HEMBA1002766 100.803 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEMBA1002769 103.210 30.236 54.098 17.099 19.753 35.636 41.922 26.940 HEMBA1002777 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776 HEMBA1002777 130.615 37.655 72.072 41.794 31.219 54.881 59.342 43.652 HEMBA1002779 97.457 29.259 75.705 22.179 22.633 33.689 38.357 77.804 HEMBA1002779 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002790 87.371 67.291 152.514 38.033 29.516 28.032 20.352 34.761 HEMBA1002801 10.311 4.603 11.704 3.190 4.420 3.016 13.823 6.693 HEMBA1002810 42.583 45.313 55.088 53.416 29.480 60.935 44.046 51.794 HEMBA1002810 42.583 45.313 55.088 53.416 29.480 60.935 44.046 51.794 HEMBA1002820 39.924 107.278 53.317 90.533 79.15 59.869 54.302 29.554 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002886 77.126 30.401 44.872 22.577 28.533 50.264 55.374 45.005 HEMBA1002886 9.474 14.188 23.688 7.557 11.980 13.85 51.233 50.264 55.374 45.005 HEMBA10022876 76.583 55.603 38.073 36.480 23.017 53.318 51.				22, 803	35, 400	15.830	15, 630	30.605	31.889	32.759
HEMBA1002755									53, 440	49, 691
HEIBRA 1002755 94.758 62.505 220.964 63.414 37.572 34.593 28.497 39.737 HEIBRA 1002759 13.935 3.117 8.450 3.792 2.291 8.714 10.261 5.285 HEIBRA 1002767 65.662 25.272 35.782 14.035 19.83 31.497 33.393 18.347 HEIBRA 1002768 100.803 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEIBRA 1002769 103.210 30.236 54.098 17.099 19.753 35.636 41.922 26.940 HEIBRA 1002777 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776 HEIBRA 1002779 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEIBRA 1002779 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEIBRA 1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEIBRA 1002794 202.405 77.515 95.182 31.252 41.834 100.167 80.301 50.036 HEIBRA 1002796 97.438 42.1334 22.488 20.281 12.823 11.156 11.567 15.735 HEIBRA 1002810 42.533 45.313 55.088 35.416 29.480 60.935 44.946 15.735 HEIBRA 1002810 42.533 45.313 55.088 35.416 29.480 60.935 44.946 17.744 HEIBRA 1002826 40.776 6.495 16.825 53.49 33.190 4.420 3.016 13.829 6.693 HEIBRA 1002826 40.776 6.495 16.825 53.49 3.319 17.745 59.865 54.302 52.958 HEIBRA 1002867 25.384 37.823 56.994 35.902 25.574 33.338 50.974 49.045 HEIBRA 100286 40.776 6.495 16.825 53.49 3.319 17.765 59.865 54.302 52.958 HEIBRA 100286 59.41 8.407 13.251 6.179 2.932 4.355 4.844 3.735 HEIBRA 100286 59.41 8.407 13.251 6.179 2.932 4.355 4.844 3.735 HEIBRA 100286 75.535 37.524 30.000 9.593 9.507 27.799 15.180 16.196 HEIBRA 100286 76.530 77.260 38.807 36.887 7.657 11.980 14.640 43.27 17.864 43.881 37.595 43.2881 43.685 43.688 7.657 11.980 43.684 35.283 42.662 42.881 37.595 43.881 37.595 43.289									20 121	
HEIBRA 1002775 13.935 3.117 8.450 3.792 2.291 8.714 10.261 5.285 HEIBRA 1002763 430.941 88.931 172.920 71.623 88.921 195.471 197.995 118.224 HEIBRA 1002767 65.682 25.272 35.782 14.035 19.183 31.497 33.393 18.347 HEIBRA 1002768 103.810 30.236 54.998 17.999 19.753 35.636 41.922 25.940 HEIBRA 1002776 103.210 30.236 54.998 17.999 19.753 35.636 41.922 25.940 HEIBRA 1002776 103.210 30.236 54.998 17.999 19.753 35.636 41.922 25.940 HEIBRA 1002777 20.350 16.768 28.054 21.736 10.754 12.030 14.991 11.776 HEIBRA 1002779 97.457 29.259 75.705 22.719 22.643 33.689 18.357 27.804 HEIBRA 1002779 97.457 29.259 75.705 22.719 22.643 33.689 18.357 27.804 HEIBRA 1002779 87.371 61.291 152.514 38.033 29.516 28.032 20.352 34.761 HEIBRA 1002794 202.405 77.515 95.182 31.252 41.834 100.167 80.301 50.036 HEIBRA 1002798 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEIBRA 10022798 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEIBRA 10022798 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEIBRA 10022810 42.583 45.313 55.088 35.416 29.480 60.935 44.946 51.794 HEIBRA 10022810 42.583 45.313 55.088 35.416 29.480 60.935 44.946 51.794 HEIBRA 1002286 52.084 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEIBRA 10022876 40.776 6.495 16.225 5.349 3.19 17.765 59.865 54.302 52.958 HEIBRA 1002286 50.735 32.524 30.030 9.593 9.527 27.595 19.197 18.101 HEIBRA 10022876 50.775 32.524 30.030 9.593 9.527 27.595 19.197 18.101 HEIBRA 1002286 75.580 27.420 49.774 44.872 22.577 28.83 50.264 55.374 45.005 HEIBRA 1002286 75.580 27.420 49.774 44.872 22.577 28.83 50.264 55.374 45.005 HEIBRA 1002286		HEMBA1002750								
HEMBA1002759 13. 935 3. 117 8. 450 3.792 2.291 8. 714 10. 261 5. 285 HEBBA1002767 65. 682 25. 272 35. 782 14. 035 19. 183 31. 497 33. 393 18. 347 HEMBA1002768 100. 803 57. 554 59. 457 35. 570 28. 006 43. 770 40. 930 38. 215 HEMBA1002769 103. 210 30. 236 54. 098 17. 099 19. 753 35. 636 41. 922 26. 940 HEMBA1002777 20. 350 16. 268 28. 054 21. 736 10. 754 12. 030 14. 991 11. 776 HEMBA1002777 130. 615 37. 655 72. 072 41. 794 31. 219 54. 881 59. 342 43. 652 HEMBA1002779 97. 457 29. 259 75. 705 22. 719 22. 643 33. 689 38. 357 27. 804 HEMBA1002790 72. 338 50. 411 181. 356 42. 070 19. 957 13. 370 27. 642 39. 672 HEMBA1002790 87. 371 61. 291 152. 514 38. 033 29. 616 28. 032 20. 352 34. 761 HEMBA1002794 202. 405 77. 515 95. 182 31. 252 41. 834 100. 167 80. 301 50. 036 HEMBA1002794 202. 405 77. 515 95. 182 31. 252 41. 834 100. 167 80. 301 50. 036 HEMBA1002796 9. 194 21. 334 22. 468 20. 281 12. 823 11. 156 11. 647 15. 735 HEMBA1002810 10. 311 4. 603 11. 704 3. 190 4. 420 3. 016 11. 829 6. 693 HEMBA1002810 42. 583 45. 313 55. 088 35. 416 29. 480 60. 935 44. 046 51. 794 HEMBA1002818 321. 516 100. 826 187. 799 84. 893 81. 585 152. 339 171. 186 117. 409 HEMBA1002820 139. 924 107. 278 533. 137 90. 533 79. 745 59. 869 54. 302 52. 958 HEMBA1002820 139. 924 107. 278 533. 137 90. 533 79. 745 59. 869 54. 302 52. 958 HEMBA1002830 119. 102 44. 248 40. 839 17. 864 23. 748 44. 398 57. 302 36. 668 HEMBA1002850 5. 941 8. 407 13. 251 6. 179 2. 932 4. 352 4. 844 3. 735 HEMBA1002867 75. 385 13. 583 42. 122 15. 283 9. 501 22. 992 15. 180 16. 196 HEMBA1002867 77. 126 30. 401 44. 872 22. 577 28. 639 50. 64 55. 374 45. 005 HEMBA1002876 101. 249 55. 603 38. 073 36. 480 23. 077 53. 318 51. 363 56. 688 HEMBA1002876 101. 249 55. 603 38. 073 36. 480 23. 077 53. 318 51. 363 56. 688 HEMBA1002876 25. 385 13. 583 42. 122 15. 283 9. 501 22. 992 15. 180 16. 196 HEMBA1002876 75. 385 13. 583 42. 122 15. 283 9. 501 22. 992 15. 180 16. 196 HEMBA1002876 37. 126 30. 401 44. 872 22. 577 28. 639 50. 64 55. 374 45. 605 HEMBA1002876 75. 385 13. 588 77. 755 774 45. 4		HEMBA 1002755	94.758	62.505	220.964	63.414	37. 572			
HEBBA1002763 430.941 88.931 172.920 71.623 88.921 195.471 197.995 118.224 HEBBA1002767 65.682 25.772 35.782 14.035 19.183 31.497 33.393 18.347 HEBBA1002768 100.803 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEBBA1002779 103.210 30.236 54.098 17.099 19.753 35.636 41.922 26.940 HEBBA1002770 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776 HEBBA1002779 97.457 29.259 75.705 22.719 22.643 33.688 59.342 43.652 HEBBA1002779 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEBBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672 HEBBA1002790 87.371 61.291 152.574 38.033 29.616 28.032 20.352 34.761 HEBBA1002798 97.457 59.582 31.252 41.834 100.167 80.301 50.036 HEBBA1002798 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEBBA1002810 42.583 45.313 55.088 35.415 29.480 60.935 44.046 51.794 HEBBA1002816 52.084 37.823 56.984 35.902 25.574 33.338 50.974 49.045 HEBBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 94.046 51.794 HEBBA1002827 139.924 107.278 533.137 90.533 79.745 59.869 54.302 52.955 8.363 HEBBA1002837 119.102 44.248 40.839 17.864 23.748 44.398 57.302 36.668 HEBBA1002866 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEBBA1002866 77.126 30.401 44.872 22.577 28.539 50.264 55.374 18.101 HEBBA1002866 9.474 14.188 23.688 7.6557 11.980 11.765 7.355 8.363 HEBBA1002866 9.474 14.188 23.688 7.6557 11.980 11.765 5.335 19.19 11.765 11.867 11.99 HEBBA1002866 9.474 14.188 23.688 7.6557 11.980 14.640 6.632 18.574 HEBBA1002866 9.474 14.188 23.688 7.6557 11.980 14.640 6.632 18.574 HEBBA1002866 9.474 14.188 23.688 7.6557 11.980 14.640 6.632 18.574 HEBBA1002876 101.249 55.603 38.073 36.480 23.077 58.881 37.595 13.890 11.801 HEBBA1002876 101.249 55.603 38.073 36.480 23.077 58.881 37.595 34.288 HEBBA1002876 101.249 55.603 38.073 36.480 23.077 58.881 37.595 34.288 HEBBA1002876 101.249 55.603 38.073 36.480 23.077 58.881 37.595 34.288 HEBBA1002876 103.2845 58.18 14.590 22.846 54.873 55.608 38.801 HEBBA1002876 103.2845 58.603 38.073 36.480 23.077 58.881 37.595 34.288 HEBBA1002874 58.500 12	30				8.450	3.792	2, 291	8.714	10.261	5. 285
HEMBA1002767 55.582 25.272 35.782 14.035 19.183 31.497 33.393 18.347 HEMBA1002768 100.803 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEMBA1002770 20.350 16.768 28.054 21.736 10.754 12.030 14.991 11.776 HEMBA1002777 130.615 37.655 72.072 41.794 31.219 54.881 59.342 43.652 HEMBA1002779 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEMBA1002779 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEMBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002790 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEMBA1002801 10.311 4.603 11.704 31.90 4.420 3.016 13.829 6.693 HEMBA1002810 42.583 45.313 55.088 35.416 29.480 60.935 44.046 51.794 HEMBA1002810 42.583 45.313 55.088 35.416 29.480 60.935 44.046 51.794 HEMBA10028270 39.924 107.278 533.137 90.533 79.745 59.869 54.302 52.958 HEMBA1002828 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002826 5.941 8.407 13.251 6.179 8.483 31.695 152.339 171.186 117.409 HEMBA1002826 50.735 32.524 30.030 9.693 9.527 27.595 19.397 18.101 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002867 25.385 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA									197 995	118 224
HEMBA1002768 100.803 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEMBA1002777 103.210 30.236 54.098 17.099 19.753 35.636 41.922 26.940 HEMBA1002777 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776 HEMBA1002777 130.615 37.655 72.072 41.794 31.219 54.881 59.342 43.652 HEMBA1002779 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEMBA1002790 87.371 61.291 152.514 38.033 29.516 28.032 20.352 34.761 HEMBA1002790 87.371 61.291 152.514 38.033 29.516 28.032 20.352 34.761 HEMBA1002794 202.405 77.515 95.182 31.252 41.834 100.167 80.301 50.036 HEMBA1002798 9.194 21.334 22.468 20.281 12.873 11.156 11.647 15.735 HEMBA1002810 42.533 45.313 55.088 35.415 29.480 60.935 44.046 51.794 HEMBA1002810 42.533 45.313 55.088 35.415 29.480 60.935 44.046 51.794 HEMBA1002810 32.156 100.826 187.799 84.893 81.685 152.339 171.186 117.409 HEMBA1002826 40.776 6.495 16.825 5.349 3.19 11.765 7.355 8.363 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002826 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBA1002826 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002866 76.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002867 25.385 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002894									33 303	
HEMBA1002779 130.615 37.655 72.072 41.794 31.957 31.370 27.642 39.672 HEMBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002799 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002799 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002799 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002798 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEMBA1002798 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEMBA1002801 10.311 4.603 11.704 3.190 4.420 3.016 13.829 6.693 HEMBA1002818 321.516 100.826 187.799 84.893 81.695 152.339 171.186 117.409 HEMBA1002818 321.516 100.826 187.799 84.893 81.695 152.339 171.186 117.409 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002837 119.102 44.288 40.839 17.864 23.748 44.398 57.302 36.668 HEMBA1002867 25.385 13.583 40.713 251 6.179 2.332 4.352 4.844 3.735 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 2.292 15.180 16.196 HEMBA1002866 7.535 33.513 583 42.122 15.283 9.501 2.292 15.180 16.196 HEMBA1002866 7.535 33.513 583 42.122 15.283 9.501 2.292 15.180 16.196 HEMBA1002866 7.535 33.513 583 42.122 15.283 9.501 2.292 15.180 16.196 HEMBA1002866 7.535 33.513 583 42.122 15.283 9.501 22.292 15.180 16.196 HEMBA1002866 7.55 335 13.583 42.122 15.283 9.501 22.092 15.180 16.196 HEMBA1002866 7.55 335 13.583 42.122 15.283 9.501 22.092 15.180 16.196 HEMBA1002866 7.55 335 13.583 42.122 15.283 9.501 22.092 15.180 16.196 HEMBA1002866 7.55 335 13.583 42.122 15.283 9.501 22.092 15.180 16.196 HEMBA1002866 7.55 335 13.583 42.122 15.283 9.501 22.092 15.180 16.196 HEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.353 56.689 HEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.353 56.689 HEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.353 56.689 HEMBA1002921 53.378 25.443 37.615 15.333 10.54 28.881 37.595 34.298 HEMBA1002931 432.841 308.291 644.522 180		HEMBA1002767	65.682	25.272	35.782	14.035			33.393	
HEMBA10027769 103. 210 30. 236 54. 098 17. 099 19. 753 35. 636 41. 922 26. 940 HEMBA10027770 20. 350 16. 268 28. 054 21. 736 10. 754 12. 030 14. 991 11. 776 HEMBA1002779 97. 457 29. 259 75. 705 22. 719 22. 643 33. 689 38. 357 27. 804 HEMBA1002779 97. 457 29. 259 75. 705 22. 719 22. 643 33. 689 38. 357 27. 804 HEMBA1002780 72. 338 50. 411 181. 356 42. 070 19. 957 31. 370 27. 642 39. 672 HEMBA1002790 87. 371 61. 291 152. 514 38. 033 29. 616 28. 032 20. 352 34. 761 HEMBA1002798 9. 194 21. 334 22. 468 20. 281 12. 823 11. 156 11. 647 15. 735 HEMBA1002798 9. 194 21. 334 22. 468 20. 281 12. 823 11. 156 11. 647 15. 735 HEMBA1002801 10. 311 4. 603 11. 704 3. 190 4. 420 3. 016 13. 829 6. 693 HEMBA1002810 42. 583 45. 313 55. 088 35. 416 29. 480 60. 935 44. 046 51. 794 HEMBA1002816 52. 084 37. 823 56. 994 35. 902 25. 574 33. 389 50. 974 49. 045 HEMBA1002818 321. 516 100. 826 187. 799 84. 893 81. 695 152. 339 171. 186 117. 409 HEMBA1002820 139. 924 107. 278 533. 137 90. 533 79. 745 59. 869 54. 302 52. 958 HEMBA1002826 40. 776 6. 495 16. 825 5. 349 3. 319 11. 765 7. 355 8. 363 HEMBA1002826 5. 941 8. 407 13. 251 6. 179 2. 932 4. 4. 392 4. 438 4. 37. 826 8. 407 13. 251 6. 179 2. 932 4. 4. 392 4. 4. 392 5. 668 HEMBA1002862 6. 941 8. 407 13. 251 6. 179 2. 932 4. 4. 392 4. 4. 392 5. 668 HEMBA1002863 77. 126 30. 401 44. 872 22. 577 28. 639 50. 264 55. 374 45. 005 HEMBA1002866 9. 474 14. 188 23. 688 7. 657 11. 980 14. 640 6. 432 18. 574 HEMBA1002886 9. 474 14. 188 23. 688 7. 657 11. 980 14. 640 6. 432 18. 574 HEMBA1002886 9. 474 14. 188 23. 688 7. 657 11. 980 14. 640 6. 432 18. 574 HEMBA1002896 78. 580 27. 420 49. 774 16. 754 20. 366 36. 684 35. 283 42. 662 HEMBA1002896 78. 580 27. 420 49. 774 16. 754 20. 366 36. 684 35. 283 42. 662 HEMBA1002921 63. 378 25. 443 37. 615 15. 333 19. 554 28. 881 37. 595 34. 298 HEMBA1002934 432. 841 308. 291 644. 522 180. 470 145. 293 27. 53. 30. 166. 153 242. 809		HEMBA 1002768	100 803	57, 554	59, 457	35.570	28.006	43.770	40.930	38.215
HEMBA1002770 20, 350 16, 268 28, 054 21, 736 10, 754 12, 030 14, 991 11, 776						17 099	19.753	35, 636	41.922	26. 940
HEMBA1002777 130.615 37.655 72.072 41.794 31.219 54.881 59.342 43.652 HEMBA1002777 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEMBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672 HEMBA1002794 202.405 77.515 95.182 31.252 41.834 100.167 80.301 50.036 HEMBA1002798 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEMBA1002801 10.311 4.603 11.704 3.190 4.420 3.016 13.829 6.693 HEMBA1002816 52.084 37.823 56.984 35.902 25.574 33.389 50.974 49.045 HEMBA1002816 52.084 37.823 56.984 35.902 25.574 33.389 50.974 49.045 HEMBA1002816 321.516 100.826 187.799 84.893 81.695 152.339 171.186 117.409 HEMBA1002820 139.924 107.278 533.137 90.533 79.745 59.869 54.302 52.958 HEMBA1002820 139.924 107.278 533.137 90.533 79.745 59.869 54.302 52.958 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002836 77.76 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002862 60.735 32.524 30.030 9.693 9.527 27.595 19.397 18.101 HEMBA1002863 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005 HEMBA1002866 75.385 13.583 42.122 15.283 50.264 55.374 45.005 HEMBA1002866 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002897 63.378 25.443 37.615 15.333 10.554 28.881 37.595 34.298 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 27.3733 166.153 242.809										
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HEMBA1002780 72. 338 50. 411 181. 356 42.070 19. 957 31. 370 27. 642 19. 672 HEMBA1002790 87. 371 61. 291 152. 514 38. 033 29. 616 28. 032 20. 352 34.761 HEMBA1002798 9. 194 21. 334 22. 468 20. 281 12. 823 11. 156 11. 647 15. 735 HEMBA1002801 10. 311 4. 603 11. 704 3. 190 4. 420 3. 016 13. 829 6. 693 HEMBA1002810 42. 533 45. 313 55. 088 35. 416 29. 480 60. 935 44. 046 51. 794 HEMBA1002816 52. 084 37. 823 56. 994 35. 902 25. 574 33. 389 50. 974 49. 045 HEMBA1002818 321. 516 100. 826 187. 799 84. 893 81. 695 152. 339 171. 186 117. 409 HEMBA1002820 139. 924 107. 278 533. 137 90. 533 79. 745 59. 869 54. 302 52. 958 HEMBA1002826 40. 776 6. 495 16. 825 5. 349 3. 319 11. 765 7. 355 8. 363 HEMBA1002826 40. 776 6. 495 16. 825 5. 349 3. 319 11. 765 7. 355 8. 363 HEMBA1002830 19. 102 44. 248 40. 839 17. 864 23. 748 44. 368 35. 57. 302 36. 668 HEMBA1002862 60. 735 32. 524 30. 030 9. 693 9. 527 27. 595 19. 397 18. 101 HEMBA1002863 77. 126 30. 401 44. 872 22. 577 28. 639 50. 264 55. 374 45. 005 HEMBA1002866 9. 474 14. 188 23. 688 7. 657 11. 980 14. 640 6. 432 18. 574 HEMBA1002886 9. 474 14. 188 23. 688 7. 657 11. 980 14. 640 6. 432 18. 574 HEMBA1002896 78. 580 27. 420 49. 774 15. 754 20. 366 36. 684 35. 283 42. 662 HEMBA1002896 78. 580 27. 420 49. 774 15. 754 20. 366 36. 684 35. 283 42. 662 HEMBA1002891 126. 001 32. 845 58. 138 14. 590 22. 846 54. 873 56. 608 38. 801 HEMBA1002896 78. 580 27. 420 49. 774 15. 754 20. 366 36. 684 35. 283 42. 662 HEMBA1002896 78. 580 27. 420 49. 774 15. 754 20. 366 36. 684 35. 283 42. 662 HEMBA1002896 78. 580 27. 420 49. 774 15. 754 20. 366 36. 684 35. 283 42. 662 HEMBA1002896 78. 580 27. 420 49. 774 15. 754 20. 366 36. 684 35. 283 42. 662 HEMBA1002921 63. 378 25. 443 37. 615 15. 333 19. 504 28. 881 37. 595 34. 298 HEMBA1002934 432. 841 308. 291 644. 522 180. 470 145. 293 273. 733 166. 153 242. 809		HEMRA1002779	97. 457	29, 259	75, 705	22.719	22.643	33.689	38.357	27.804
HEMBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEMBA1002794 202.405 77.515 95.182 31.252 41.834 100.167 80.301 50.036 HEMBA1002798 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEMBA1002810 10.311 4.603 11.704 3.190 4.420 3.016 13.829 6.693 HEMBA1002810 42.583 45.313 55.088 35.416 29.480 60.935 44.046 51.794 HEMBA1002818 321.516 100.826 187.799 84.893 81.695 152.339 171.186 117.409 HEMBA1002810 331.516 100.826 187.799 84.893 81.695 152.339 171.186 117.409 HEMBA1002820 139.924 107.278 533.137 90.533 79.745 59.869 54.302 52.958 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002833 119.102 44.248 40.839 17.864 23.748 44.398 57.302 36.668 HEMBA1002867 5.941 8.407 13.251 6.179 9.527 27.595 19.397 18.101 HEMBA1002867 5.941 8.407 13.251 6.179 9.527 27.595 19.397 18.101 HEMBA1002867 75.385 13.583 42.122 15.283 9.501 22.392 15.180 16.196 HEMBA1002886 9.474 14.188 23.688 7.557 11.980 14.640 6.432 18.574 HEMBA1002886 9.474 14.188 23.688 7.557 11.980 14.640 6.432 18.574 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.664 35.283 42.662 HEMBA1002891 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEMBA1002891 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEMBA1002921 53.378 25.443 37.615 15.331 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809						42 070	19 957	31, 370	27, 642	19,672
HEMBA1002794 202.405 77.515 95.182 31.252 41.834 100.167 80.301 50.036 HEMBA1002798 9.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735 HEMBA1002810 10.311 4.603 11.704 3.190 4.420 3.016 13.829 6.693 HEMBA1002810 42.583 45.313 55.088 35.416 29.480 60.935 44.046 51.794 HEMBA1002816 52.084 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEMBA1002818 321.516 100.826 187.799 84.893 81.695 152.339 171.186 117.409 HEMBA1002820 139.924 107.278 533.137 90.533 79.745 59.869 54.302 52.958 HEMBA1002820 40.776 6.495 16.825 5.349 31.11 17.65 7.355 8.363 HEMBA1002826 40.776 6.495 16.825 5.349 31.11 17.65 7.355 8.363 HEMBA1002833 119.102 44.248 40.839 17.864 23.748 44.198 57.302 36.668 HEMBA1002862 60.735 32.524 30.030 9.593 9.527 27.595 19.397 18.101 HEMBA1002862 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005 HEMBA1002876 101.249 55.603 38.073 16.480 23.017 53.318 51.363 56.689 HEMBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 37.595 34.298 HEMBA1002894 432.841 308.291 644.522 180.470 145.293 27.3733 166.153 242.809 432.841 432.841 4308.291 644.522 180.470 145.293 27.3733 166.153 242.809 432.841 432.841 4308.291 644.522 180.470 145.293 27.3733 166.153 242.809 432.841 432.841 4308.291 644.522 180.470 145.293 27.3733 166.153 242.809			12.336							
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HEMBA1002801 10.311 4.603 11.704 3.190 4.420 3.016 13.829 6.693 HEMBA1002810 42.583 45.313 55.088 35.416 29.480 60.935 44.046 51.794 HEMBA1002816 52.084 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEMBA1002818 321.516 100.826 187.799 84.893 81.695 152.339 171.186 117.409 HEMBA1002820 139.924 107.278 533.137 90.533 79.745 59.869 54.302 52.958 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002830 19.102 44.248 40.839 17.864 23.748 44.398 57.302 36.668 HEMBA1002850 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBA1002862 60.735 32.524 30.030 9.693 9.527 27.595 19.397 18.101 HEMBA1002863 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEMBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002896 78.580 27.420 49.774 15.754 20.366 36.684 35.283 42.662 HEMBA1002891 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809		HEMBA1002798	9, 194	21, 334	22, 468	20. 281	12.823	11, 156	11.547	15.735
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HEMBA1002816 52.084 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEMBA1002818 321.516 100.826 187.799 84.893 81.695 152.339 171.186 117.409 HEMBA1002820 139.924 107.278 533.137 90.533 79.745 59.869 54.302 52.958 HEMBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBA1002833 119.102 44.248 40.839 17.864 23.748 44.198 57.302 36.668 HEMBA1002850 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBA1002862 60.735 32.524 30.030 9.693 9.527 27.595 19.397 18.101 HEMBA1002863 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HEMBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002921 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809										
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HEMBA1002860	40	HEMBA 1002833	119.102	44. 248	40.839	17.854	23.748			
HEMBA1002862 60.735 32.524 30.030 9.693 9.527 27.595 19.397 18.101 HEMBA1002863 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HEMBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002913 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEMBA1002921 53.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809				8, 407	13, 251	6, 179	2.932	4. 352	4.844	3.735
FEMBA1002863 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005 FEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 FEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 FEMBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 FEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 FEMBA1002813 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 FEMBA1002921 53.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 FEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 FEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809										
HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HEMBA1002886 9.474 14.188 23.688 7.557 11.980 14.640 6.432 18.574 HEMBA1002891 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002913 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEMBA1002921 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809 55								FA 364		
50 HEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HEMBA1002886 9.474 14.188 23.688 7.557 11.980 14.640 6.432 18.574 HEMBA1002896 76.560 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002891 126.001 32.845 58.138 14.590 28.86 54.873 56.608 38.801 HEMBA1002921 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809		[HEMBA1002863	77.126	30.401	44.872					
50 HEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HEMBA1002886 9.474 14.188 23.688 7.557 11.380 14.640 6.432 18.574 14.640 6.432 18.574 14.640 6.432 18.574 14.640 6.432 18.574 14.640 6.432 18.574 14.640 6.432 18.574 16.754 20.366 36.684 35.283 42.662 14.640 14		HEMRA 1002867		13, 583	42, 122	15, 283	9. 501	22.992	15.180	16.196
FINAL PROPERTY OF THE PROPERTY							23 017	53.318	51, 363	56, 589
HEMBA1002886 9,4/4 14,188 23,686 7,637 11,380 42,530 42,662 HEMBA1002896 78,580 27,420 49,774 16,754 20,366 36,664 35,283 42,662 HEMBA1002813 126,001 32,845 58,138 14,590 22,846 54,873 56,608 38,801 HEMBA1002921 63,378 25,443 37,615 15,333 19,054 28,881 37,595 34,298 HEMBA1002924 65,007 29,109 104,125 15,411 19,920 31,099 23,998 19,182 HEMBA1002934 432,841 308,291 644,522 180,470 145,293 273,733 166,153 242,809 55	50									
HEMBA1002913 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEMBA1002921 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809		HEMBA 1002886								
HEMBA1002913 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEMBA1002921 53.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809		HEMBA1002896	78.580	27. 420	49.774	16.754				
HEMBA1002921 53, 378 25, 443 37, 615 15, 333 19,054 28,881 37,595 34,298 HEMBA1002924 65,007 29,109 104,125 15,411 19,920 31,099 23,998 19,182 HEMBA1002934 432,841 308,291 644,522 180,470 145,293 273,733 166,153 242,809						14.590	22.846	54.873	56.608	38.801
HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809										
55 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809										
55 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809		HEMBA1002924	65.007	29.109						
55				308, 291	644.522	180.470	145. 293	273.733	166.153	242.809
UEMONIANCE 22 25' AND 1 ATT THE TEST 41' AND 1 ATT DATE 25' AND 1 ATT DATE	55								29.646	38, 889
		UCHON LONGAS	92.005	32.104	1 441.144	1 73.711	77.007	1 54.55.		

	HEMBA1002937 T	20 600 1	30, 844	33,817	12.784	18. 251	14, 107	24. 131	18.662
		38. 698	22.867	33.838	19.077	13.734	19. 266	17.364	17.750
	HEMBA 1002939	39.755			21.860	18. 241	23. 920	21, 112	16. 286
5	HEMBA 1002944	53, 752	33.349	51.861		29.614	19.702	28. 422	21.177
Ū	HEMBA 1002951	38. 716	29.783	39.196	19.808				13. 424
	HEMBA 1002954	24. 907	8. 542	20.941	9. 265	13.758	15.056	7.297	
	HEMBA 1002962	86.680	62.578	220. 246	62.027	37.753	44.037	31.812	41.725
	HEMBA 1002968	105. 871	78,850	221.414	65, 545	40, 380	43.093	38.816	50. 281
	HEMBA 1002970	48. 034	34,741	30.834	18.482	6.639	17. 125	23. 514	36. 180
10	HEMBA1002971	39, 492	44.145	35.618	25.614	12. 932	25. 193	14.823	23. 202
10	HEMBA1002973	83.710	70.965	156, 167	43.307	28. 902	29, 947	26. 101	34. 769
	HEMBA1002978	35. B33	19. 362	27.056	13.075	20. 398	11.324	16.059	13. 956
	HEMBA1002981	107. 112	35, 200	56.576	23.895	26.105	33.054	17.199	21.249
	HEMBA1002985	79. 217	44. 154	116,532	27.950	26.158	37.452	28. 927	20. 335
	HEMBA 1002986	61, 056	78.203	68.834	49.967	64. 529	38. 333	28.919	20. 529
45	HEMBA 1002988	37, 307	36.609	71,802	20.621	8.965	16.229	15. 956	22.796
15	HENBA1002992	97.720	72.656	79.841	50. 454	34. 289	57.004	61.291	91.211
	HEMBA1002995	51.473	63.779	55, 081	36.903	25.007	38.630	19.510	48. 529
	HEMBA1002997	41.734	70.805	29.264	27.019	33.664	24. 201	18.442	25.973
	HENBA1002999	35. 341	16.456	18. 357	11.146	7,034	12.086	13.966	9. 970
	HEMBA 1003 004	55. 654	33.689	35. 194	15.119	16. 204	20.866	27.891	20.055
	HEMBA 1003006	40. 682	24.886	20.750	20.903	26.595	25. 445	20.310	20. 924
20	HEMBA 1003008	29. 269	20, 922	74.697	25.061	17.787	10.271	5.688	12.638
	HEMBA 1003003	130, 889	123.646	311.225	101.957	95. 443	64.844	60.969	90. 296
	HEMBA1003027	54. 935	32.610	44.710	18.890	52. 131	26. 286	28.112	31.561
	HEMBA 1003029	33. 333	42.436	60. 787	20.829	34.111	29.704	49.230	45. 833
	HEMBA 1003029	34, 000	25.311	18. 494	14. 998	13.316	13.955	15.773	27. 136
	HEMBA 1003032	171.114	46.990	71, 365	23.640	50.526	81.278	84.036	46. 352
25		168. 563	118.674	378.771	109. 222	90.670	70. 150	55. 336	77.819
	HEMBA 1003033 HEMBA 1003034	173. 162	127.221	484. 135	108. 218	85.630	61.733	36.799	63. 312
	HEMBA 1003035	11.693	5. 195	9.305	4. 478	5. 058	11.024	2.553	4. 409
	HEMBA 1003037	261.159	89.481	145, 321	58. 521	65.732	104.677	89, 571	71.674
	HEMBA1003041	103. 945	105.085	291.931	93.188	75. 193	53.097	39.564	58. 217
	HEMBA 1003041	40. 254	39, 965	46.856	26.192	11.615	35. 659	25.378	32.416
30	HEMBA1003047	127. 888	49. 341	139.750	32.219	32.320	57. 450	33.390	28.702
	HEMBA 1003048	87. 433	35. 962	42.305	12.040	20, 442	39, 108	29.597	21.461
	HEMBA1003064	6. 366	8. 535	6. 201	8.809	4. 415	7. 239	3.330	7.829
	HEMBA1003067	55. 833	34.508	77.097	26. 154	20.523	28.755	24.783	17.488
	HEMBA1003071	54. 728	22.509	28. 869	17.461	19.647	20.624	22.285	19.438
	HEMBA1003072	62. 421	30.769	31.225	26.146	22.906	21.483	17.616	19.134
35	HEMBA1003076	111. 254	51.085	78. 972	37.151	40.422	49.911	47.023	64.737
	HEMBA1003077	36.471	15. 407	24. 522	8.009	8, 453	18.661	13.797	5. 837
	HEMBA1003078	34, 143	38.741	77.906	31.907	37.169	17.933	17.439	18. 923
	HEMBA1003079	28. 559	39.563	41.646	26.110	25.889	25.576	18.026	24. 526
	HEMBA1003083	61.036	48.635	169.439	52.788	60.016	41.611	29.619	67.469
	HEMBA1003086	49. 032	40.488	154, 409	29.869	12.063	16.544	16.039	19.219
40	HEMBA1003090	34.778	14.860	23.758	12.710	24.132	15.848	25.027	14. 265
	HEMBA1003094	184. 999	43.363	72.116	30.096	53.636	78.251	84.551	34.775
	HEMBA 1003096	31. 440	18.030	25.774	10.290	11.781	14.033	27.791	11.348
	HEMBA1003098	36, 774	64.970	88.562	34.074	24.271	25.656	18.003	31.059
	HEMBA 1003101	55.716	24.121	22.316	11.682	13.163	21.315	25.117	15. 689
4=	HEMBA1003109	48. 411	21.093	39, 285	21.315	21.724	27.826	31.034	21.809
45	HEMBA1003114	41, 101	24.785	22.792	14.164	14.657	18. 320	15.152	16.038
	HEMBA1003117	22. 939	13.535	20, 191	6.812	10.538	14.917	18.015	12.566
	HEMBA1003120	24. 531	24.408	55.805	26. 574	13.838	15. 423	15.080	21,728
	HEMBA1003129	40. 276	46.792	104.463	37.995	37, 989	21.990	26.267	18. 207
•	HEMBA1003133	50.080	22.873	35.022	15.164	20.000	21.592	25.551	27. 656
EO	HENBA1003136	146, 630	23.706	65.990	18.301	31.049	69.754	51.669	25. 346
50	HEMBA1003142	69.008	47.867	130.557	32.955	30.384	25. 274	27.118	29. 493
	HEMBA 1003 148	59. 282	20.084	32.740	18. 292	18.973	32. 206	22.003	24.674
	HENBA1003151	53. 856	20.003	51.824	13, 233	9.854	27.114	22.251	13, 546
	HEMBA1003152	20. 577	9. 803	19. 388	10.017	5.761	31.586	23.227	6.853
	HEMBA1003157	16. 477	9. 272		9.919	17.605	7.547	10.156	10. 181
66	HEMBA1003166	293. 814	257. 380		260. 521	221.325	137. 459	148.208	199.758
55	HEMBA1003171	17. 730	8.702		6, 499	6.963	7.361	5.733	7.164
	(UCHON 1003111	1 11.130	3. 102		-1	1			

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	FIRM LANGUAGE T		10 (16 1	400 100 T	30 FOA I	17.624	21.152	13.386	15. 936
	HEMBA1003175	38.620			29. 594				
	HEMBA1003179	63.835	33.869	50.631	27.163	25. 502	35.500	39.052	37.713
	HEMBA1003186	100, 461	75.611	231.787	75.781	58.278	54.222	55.862	61.615
5					20. 523	18.740	21.756	30. 501	35. 864
	HEMBA1003196	36.422	27.557	45. 633					
	HEMBA1003197	8. 462	9.564	5. 534	5, 965	4. 051	3.138	7.054	7.066
	HEMBA1003199	34, 650	18, 409	81. 183	15.696	16.799	9, 492	17.381	15. 917
	HEMBA1003202	79. 337	59.764	236.822	43.286	41.820	31.106	32, 936	45. 183
						31, 451	33.193	27.421	28. 849
	HEMBA1003204	66. 523	56. 272	172.818	48.560				
10	HEMBA1003210	23.713	52.768	35.498	5. 529	38.451	16.353	59.417	17.563
10	HEMBA1003212	126. 394	90.709	372.474	74.164	52. 392	59.663	45_714	54. 363
		19. 415	13, 105	13.670	6.371	4.792	13.681	10.789	6. 536
	HEMBA1003218					42. 391	47.586	54.647	123.019
	HEMBA1003220	81, 171	86.642	147.453	89.495				
	HEMBA1003222	25. 803	22.891	28. 577	7.994	11, 404	10.413	9.856	14. 985
	HEMBA1003225	105, 735	21, 238	40.848	11.586	20.280	48.243	44.574	19. 547
	HEMBA1003229	30.394	26. 363	41.333	22.998	17.475	14.707	20. 154	19.749
15						20, 775	56.815	40, 191	75. 238
	HEMBA1003230	69.643	70.015	42.439	31.176				
	HEMBA1003235	44.989	43, 337	105. 267	33.038	19. 405	20.834	22.018	29. 856
	HENBA1003236	8.677	17.896	8.735	7.270	7. 328	17.286	5. 295	18.441
	HENBA1003250	7.260	12.5981	12.993	4.750	4.815	7.242	5. 982	4, 378
		65 274		65. 197	28.241	33, 512	44.917	62.506	60, 076
	HEMBA1003252	56.274	51.495						
20	HEMBA1003257	71.751	16.083	40.414	13.391	19.441	38.988	28.614	19.028
	HEMBA1003268	19.492	18.996	46. 948	14.167	12.769	11.524	8. 622	17.414
	HEMBA1003273	48.113	38.933	125, 242	29.404	21.135	22.989	17.240	24. 704
	HEMBA1003276	36.279	34. 802	113.584	23.812	17. 208	20. 437	14.685	26. 145
						11. 287	13.206	16. 182	14. 465
	HEMBA1003277	31.353	12.827	21.514	10.462			10. 948	17. 841
	HEMBA1003278	36.998	24. 906	71.222	17.479	15.791	16.787		
25	HEMBA1003280	50.716	16.000	38. 057	16.933	20.792	37.901	30.931	31.493
23	HEMBA1003281	66,732	21. 393	32.728	15.032	18.415	26.844	28.577	24.898
		9.746	8. 482	12. 941	5.779	5. 747	5.813	3.545	3.499
	HEMBA1003284					29, 473	52.233	50. 283	47.695
	HEMBA1003286	69.502	35. 947	60.729	21.827				
	HEMBA1003291	13.248	9, 951	10.909	3. 504	18. 100	6. 561	6. 341	7_647
•	HEMBA1003294	69.599	52, 239	168. 555	39. 127	38. 460	40.377	24.057	27. 486
	HEMBA1003296	61.933	31.456	37.947	21,206	23. 199	23.249	34, 580	37.768
<i>30</i>			5. 972	8.976	6.154	8.839	4, 199	3, 451	3. 227
	HEMBA1003304	7.117					8.448	6. 333	9. 387
	HEMBA1003306	17. 590	15. 590	22.443	8.410	11. 282			
	HEMBA 1003309	5.845	10. 103	12. 198	14.015	7.776	8,709	3, 955	18. 326
	HEMBA1003314	637.052	210.608	238.618	105.098	198. 106	299, 884	273.738	171, 516
4	HEMBA1003315	83.736	51,612	84.690	32. 381	29, 482	56.694	53, 105	54. 024
				256.570	51.502	51.083	44, 130	42.804	45. 519
35	HEMBA1003322	108. 401	88.539				18, 087	12.420	9. 516
	HEMBA1003326	42, 723	20.581	14.759	11.799	7.780			
	HEMBA1003327	61.811	36.702	87.698	28. 181	19, 784	18.596	17.453	18. 377
	HEMBA1003328	53. 406	51.712	114.941	36.926	25.000	18.669	22.079	32.865
	HEMBA1003330	108. 955	82.099	207.708	73.413	52.244	50.838	55.920	55, 390
			110. 275	337. 182	94. 209	99.717	67.000	43.513	80.023
	HEMBA1003348	121.625					9.064	3, 020	5, 373
40	HEMBA 1003369	5.861	23.644	14.930	4.979	1.726			
	HENBA1003370	315.016	197.956	369.117	140.044	139.216	140.758	150. 458	124, 948
	HEMBA1003373	50.135	31.291	53.330	17, 430	5.513	19.164	8.117	19, 638
	HEMBA1003376	174. 269	170.290	519.668	125.099	89, 798	108.226	81.818	107.084
		43.015	24.657	74.071	29.281	24, 407	19,711	13, 485	20.047
	HEMBA1003380			68.079	15. 389	9. 455	11.810	8.800	14. 281
	HEMBA1003384	25, 555	30.071						
45	HEMBA 1003387	6. 515	2.588	2.697	1.577	1. 109	1.803	1.986	3.464
73	HEMBA1003392	111. 457	25.882	42. 253	17. 323	29.007	50,086	29.337	23. 550
	HENBA1003395	16.068	18.666	35. 483	15.254	9.873	10.355	6. 207	12.514
			21.480	37.035	19.231	15. 354	19.471	27.860	34, 116
	HEMBA1003399	45. 227				49, 133	53.819	60.041	53.109
	HENBA 1003400	116. 210	36.907	58.706	24.811				
	HEMBA1003402	32.500	16.239	27.864	8.795	12.857	17.141	11.617	14, 596
50	HENBA 1003 403	60.260	43, 377	46.720	20.221	26.579	36.738	44.891	45, 870
50	HENBA1003408	196. 576	49.687	70.460	29.354	50, 910	84. 358	77.062	46.433
				55. 699	47.250	43.763	61,953	59, 463	47. 139
	HEMBA1003412	104, 813	43. 934						11.032
	HEMBA1003417	22. 445	13.970	25. 036	8.433	7. 282	10.593	5.696	
	HEMBA1003418	57.411	57. 397	76.232	97.795	45. 336	43, 450	22.206	90. 604
	HEMBA1003420	29.838	15.856	201.831	11.319	8.067	11.379	12.938	14, 721
	HEMBA1003425	17. 466	15. 895	21. 662	4.733	6, 723	8.483	10, 838	9.083
55						10. 421	10.074	11.092	11.581
	HEMBA1003433	23, 931	18. 435	24. 576	12.136	1 10.461	1 10.074	11.032	1 401

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	HEMBA1003440	91.727	41.727	39. 257	19.755	26.941	45. 998	31.620	35. 845
	HEMBA1003442	7.090	22.535	10.452	33. 897	10.259	15.118	7.093	14.790
	HEMBA1003447	82, 161	36.670	48.248	26.789	18.587	41.591	42.314	35.065
5	HEMBA1003453	50. 472	25.692	25. 954	16.130	11.252	16.584	28. 534	21. 256
	HEMBA1003461	55. 687	25. 328	42.686	17. 261	18.856	27. 281	22.795	17. 854
	HEMBA1003463	40, 102	23, 311	34.469	13. 456	19.704	20, 277	16.984	18.124
	HEMBA1003465	92. 245	40.963	61.816	28.410	36.051	39, 389	40. 220	36.851
	HEMBA1003480	114.075	114, 841	256.076	76. 366	67.942	56.459	51, 589	62. 191
	HEMBA1003485	44. 403	28.836	33.659	14. 371	8.636	26. 284	16.036	14. 582
10	HEMBA1003487	42.939	15. 463	23.730	9.752	15.729	24. 902	21.136	16.494
		31.026	21.538	56.674	14. 934	12.014	12.082	9.567	14.655
	HEMBA1003492	97.366	260. 496	50, 174	48. 821	12.504	74. 554	20.623	180.841
	HEMBA1003494		17.943	24. 659	11, 432	13.881	21.376	18. 562	6.072
	HEMBA1003497	19.000	21. 486	28. 175	12. 948	17.154	30.911	36.463	16.806
	HEMBA1003503	54.774 18.672	14.740	43.023	11, 794	13.330	8. 925	16. 405	11.615
15	HEMBA1003511	385. 123	191. 234	239.319	81. 329	123.915	213.945	179.430	96.672
	HEMBA 1003528		12.384	23.693	10, 695	21.216	20.067	28.030	16.204
	HEMBA 1003530	43.820	73.542	215.578	67, 833	214.022	56. 139	50. 217	66.992
	HENBA1003531	111.104	62.379,	83.827	37. 506	53. 388	90. 314	77.728	60.515
	HEMBA 1003532		20.746	32.949	11, 160	19. 286	34, 305	28. 231	13.837
	HEMBA 1003538	61.123	10.501	20,608	5. 904	7, 197	10. 239	6.617	8.168
20	HEMBA 1003545 HEMBA 1003546	21.489 31.371	32.365	28.613	13. 365	225.243	16. 427	16.554	24. 821
	HEMBA1003548	4, 456	8. 124	9. 845	4. 563	7.542	6.155	5.647	8.387
	HEMBA1003553	79.837	51.515	50. 379	23. 127	28.564	49, 154	63. 525	48. 955
	HEMBA 1003555	20.066	8. 873	13.692	4. 762	3.684	10.112	10.962	6. 521
		57. 280	36.399	128. 391	29. 283	16, 426	19. 257	18. 121	24.622
	HEMBA1003556 HEMBA1003560	9. 290	4. 426	2.529	2.848	1,767	2.983	6. 207	6, 539
25	HEMBA 1003565	42.648	29. 588	20. 996	8.344	13.984	21. 927	21.847	22.043
	HEMBA 1003568	7. 244	1.649	7.712	2. 430	3.763	3. 172	2. 836	2, 592
	HEMBA 1003569	25, 048	20. 536	23.764	33.957	13.740	16.235	19.512	16, 518
	HEMBA 1003571	111.721	94. 378	126.335	84, 368	71.788	50.029	48.011	59, 960
	HEMBA 1003579	3.335	7. 399	15.353	6. 553	8.948	2.872	9. 198	6. 421
	HEMBA 1003580	274. 105	50. 292	102.103	26.686	59.875	128. 943	110.375	35, 695
30	HEMBA 1003581	112.013	31. 295	94. 083	21.641	36.215	54. 336	50.711	21.238
	HEMBA 1003591	97.076	64. 326	77. 160	89.876	47.882	53.615	40.656	45.172
	HEMBA 1003595	32.697	22.842	84.629	19.075	11,339	6. 305	5. 581	18.085
	HEMBA 1003597	48, 561	25.846	108.491	20.931	15.952	19.375	17.580	20, 153
	HEM8A1003598	49.728	20.134	22.468	12.142	11.688	18.934	21.743	15.025
	HEMBA 1003600	32.772	35.099	56.905	26.268	29.290	38.873	53. 305	56.783
35	HEMBA1003602	18. 248	10.116	16.162	6.182	10.970	8.064	14.736	17.188
	HEMBA1003604	205. 949	53.579	69.723	24.549	49.902	105. 181	98, 166	47. 144
	HEMBA 1003610	140.996	29. 255	95.048	15.492	103.150	72.233	54.670	30,688
	HEMBA 1003615	57.258	20.035	34, 102	12.808	16.022	24. 378	18.759	20.876
	HEMBA1003617	48.414	20.375	29, 789	12.148	22. 291	18. 199	18.770	18. 242
40	HEMBA1003620	52.899	22.318	45.502	19.575	19.962	25. 239	39.072	29. 451
40	HEMBA1003621	102.827	102.094	226.373	80.194	64.742	58.874	67.142	60.680
	HEMBA1003622	19.815	13.838	25.009	16.055	8. 339	12. 261	15. 369	13.833
	HEM8A1003630	20.008	16.381	30.244	13.871	5. 573	9. 992	10.303	11.422
	HEMBA 1003637	37.880	29.848	106.379	23.251	18.468	17. 181	12.409	18.500
	HEMBA 1003640	39.068	31.672	100, 901	22.572	22. 223	21.513	17.417	20. 420
45	HEMBA 1003645	25.820	19, 380	48. 445	13.481	9, 247	12. 142	54. 230	5,711
40	HEMBA1003645	38.243	15. 329	22.003	9.624	13.311	24,606	19, 177	19.938
	HEMBA 1003647	10.261	10.718	12.323	7.860	7.892	7.607	7.882	10.058
	HEMBA 1003656	40.171	31.269	66.874	28. 981	19, 429	18.898	23.172	30.178
	HEMBA 1003662	25.325	17.011	19.352	6.387	10.041	10.909	14.055	18.544
	HEMBA 1003666	23.085	11. 187	17.407	5.803	8. 262	9.774	15.332	13.851
50	HEMBA 1003667	304.975	209. 929	337.134	96.636	131.792	179, 317	140.769	174.256
	HEMBA 1003670	12.944	8.894	15. 235	3.344	2. 565	7.057	6. 425	7.073
	HEMBA1003674	143.262	32, 196	51.919	33.863	62.734	66.675	65, 424	47.173
	HEMBA1003677	80.516	45. 946	220.695	45.985	43.474	38, 916	30. 594	46.808
	HEMBA 1003679	25.325	7.795	16. 167	6.727	5. 941	12. 433	12.034	11.720
	HEMBA1003680	42,317	25.723	33, 794	24.664	23.985	25. 419	38.990	39.343
55	HEMBA 1003684	18.273	10.175	17.733	13. 315	4. 937	9. 099	10.182	10.574 56.592
	HEMBA 1003690	115.021	65, 531	75.876	46. 124	43.039	71.797	85. 431	1 30.334

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								40 000	40.544
	HEMBA1003692	83. 253	96.347	194. 372	63. 188	40.872	47. 354	39. 288	42.644
	HEMBA1003702	88. 125	35.028	48. 251	23.719	29.023	42.879	46.956	36. 550
_	HEMBA1003711	93. 732	50.280	140. 199	32.885	33.424	47. 500	41, 959	36. 807
5	HEMBA1003714	75. 923	20.696	37.340	14. 414	28.237	32.029	29, 145	16. 214
	HEMBA1003715	54. 160	54.486	142.871	31.894	31.122	28. 832	20, 640	26.672
	HEMBA1003717	70.553	38, 574	120.922	45. 101	29. 491	29. 344	27, 200	38. 418
	HEMBA1003720	83. 587	94.829	133.285	55. 896	49.519	43. 330	22.099	41, 137
	HEMBA1003725	46. 157	55. 932	71.704	30.085	21. 305	22. 378	18.643	31.573
	HEMBA1003728	103.795	35.658	58. 184	16.485	21.818	42. 286	37.790	34. 280
10	HEMBA1003729	49. 957	21.508	47.663	20.231	15.376	18. 567	21.294	17. 427
	HEMBA1003732	13.069	1, 953	6, 558	3. 228	2.195	3.652	3.024	4. 336
	HEMBA1003733	52.409	32.781	76.684	22, 919	83.426	18.921	13.867	14. 220
	HEMBA1003742	40. 426	20. 265	50.667	26. 589	21.518	42.057	44.130	24. 802
	HEMBA 1003743	26. 918	22.118	23. 392	18.886	18.530	12.506	17.162	18.069
	HEMBA1003758	110.630	126.359	315.104	79. 435	58.130	58. 587	34.868	73. 429
15		78. 949	0.000	26.318	15, 194	14. 440	32.057	34.468	19. 471
	HEMBA1003760	45. 855	30.390	82.720	23. 891	19.630	184. 051	37.797	57. 861
	HEMBA1003764	87.589	47. 227	62. 942	27. 144	32.047	46, 499	39.296	38. 944
	HEMBA1003769			21, 132	12,002	9.850	33.904	29, 817	13. 165
	HEMBA1003773	63.842 17.751	14.722 16.975	23.942	16.465	13.884	6. 842	9. 757	20. 650
	HEMBA1003783		17 222		13.856	12.436	17. 394	11.099	13.140
20	HEMBA 1003784	13.500 386.642	17, 233 303, 008	21.849 322,299	109.371	145.316	286.778	287.377	239. 938
	HEMBA1003794			29.603	15.022	13.775	16. 550	24.428	19. 403
	HEMBA 1003799	39, 392	23.099 21.899	44. 323	20.132	18.580	28. 795	24.744	35. 938
	HEMBA1003803	63. 548		44.323	16.154	27.867	31.087	37.611	22. 634
	HEMBA 1003804	80.382	26.816	48.558	19. 994	36.377	43.797	32.147	28. 376
	HEMBA 1003805	103.669	42.485	42.930	9, 492	6.870	9. 649	8.812	7.611
25	HEMBA 1003807	21.717	13. 940	25. 512		17.855	7. 640	4. 451	6. 585
	HEMBA 1003810	20, 102	11.572	7. 558	20. 338 155. 416	219.584	266.037	283.204	241. 127
	HEMBA 1003827	432. 964	219.520				93, 790	92.728	122. 237
	HEMBA 1003836	177. 311	135.831	482. 134 641. 368	146.466	136.063 79.993	115, 711	87. 137	118.957
	HEMBA1003838	223. 674	185. 295		17.850	21. 104	13. 382	11.701	13.634
	HEMBA1003843	13.867	10.178	27. 409	34. 962	50.550	56. 395	40.851	60.253
30	HEM8A1003846	133.994	57. 556	58.738	14. 248	8.662	11. 259	9. 145	9.934
00	HEMBA1003856	27. 378	13.868	16. 982		52.628	51. 958	45. 837	48.871
	HEMBA1003857	101.908	95. 527	253. 525	75, 110	13.009	16.810	29.271	16.795
	HEMBA1003864	52.130	18.071	24. 567	9, 568	15.414	19. 103	9. 229	7.524
	HEMBA1003866	27.257	12.805	22. 440	12.069 31.090	41.733	69. 461	48.174	43.486
	HEMBA1003868	95.701	54. 991	58. 923		42.446	37. 097	36.010	45. 824
35	HEMBA 1003879	62.950	44.572	159. 217	48. 098	47. 956	67.668	44. 498	30.581
50	HEMBA 1003880	134, 462	70, 074	103. 271	50.699 34.796	54.399	57. 269	63.551	68.830
	HEMBA1003884	99, 190	48. 465	73.499	55, 129	49.424	41.309	24. 247	31.596
	HEMBA1003885	77.675	69.096	172.968 33.582	16.896	21.181	29. 281	31.275	22.835
	HEMBA1003887	60. 203	22.185		7.762	16.057	139, 271	387, 408	5.124
	HEMBA 1003890	12.753	8.056	15.506	187.300	180.355	212. 964	137. 297	122. 335
40	HEMBA1003893	386. 525	281.955	515, 307 382, 182	144.104	165. 806	233. 857	186.700	143.577
70	HEMBA1003896	411.418	232.899 39.491	114. 984	20, 297	23. 509	16.793	14. 124	20.479
	HEMBA 1003902	39.732 32.775	21.109	45.629	10.006	13.109	14. 294	24.342	17.444
	HEMBA1003904	8.660	8. 873	15.689	7.298	15. 429	6.307	2.267	5.699
	HEMBA1003908		253.614	316.882	183.017	124. 195	147. 955	105, 962	360.995
	HEMBA1003926	132.636	63.862	200.940	40.687	36.238	35. 284	29.695	40.418
45	HEMBA 1003937 HEMBA 1003939	87.005 28.064	25, 844	35.675	20, 306	20.378	19.070	16. 457	15.626
40	HEMBA1003940	27, 800	13.368	18.045	10. 235	10.394	14.633	17,733	9.868
			16.835	24.582	17. 381	15.884	23.428	19, 757	13,795
	HEMBA1003941	57.997	19.747	45. 852	32.660	22.333	24.695	10.791	21,900
	HEMBA1003942	38.168	32. 900	46.079	23.037	21.163	36, 632	32. 279	26. 903
	HEMBA1003945	59.457	13.019	20.678	7.159	38. 521	442. 120	272.494	21.625
50	HEMBA 1003949	12.870	8.726	5.814	3, 195	4. 756	3.396	8.814	5. 401
00	HEMBA 1003950	8.366	10.310		9, 390	8. 494	10.637	10.973	5. 252
	HEMBA 1003953	23.527	90.718	253.084	74, 499	85.036	62. 450	34. 852	86.629
	HEMBA1003958	131.082	11. 228	18.520	6. 548	7, 960	18. 122	12.612	10.591
	HEMBA1003959	12, 105	29.785		18, 932	16. 178	21. 708	32.094	35, 333
	HEMBA1003960	53.133 58.245	19.415		20.791	26. 975	28. 975	27.825	25.303
55	HEMBA1003966	1.859	3. 908			4.054	4. 384	4. 208	4.986
	HEMBA1003967	1. 639	3. 300	3. 384	1 0.000	7.034	1		

	DEMESTARANCO T	40 310 [40 004	CE 257	16.296	14, 511	28, 531	22. 648	15, 420
	HEMBA1003968	40.219	26.894	55. 357		33. 973		29. 356	338.820
	HEMBA1003974	147. 167	419.547	139.030	117.010		54. 122		
5	HEMBA1003976	20.167	17. 809	13.159	9. 187	5.748	6.820	6. 962	10. 167
•	HEMBA1003977	32.761	12. 350	24.212	6. 558	6,776	12.413	17.016	9.367
	HEMBA1003978	40. 564	13. 858	10.812	11.585	11. 203	23, 881	20. 489	17.488
	HEMBA1003981	65. 803	34. 462	71.399	26.801	31. 348	48.051	31. 355	42.728
	HEMBA1003982	15, 104	89.360	20. 946	18.086	1.620	3. 781	3. 102	64.356
	HEMBA1003985	15, 199	10.866	21.715	9. 199	1.517	8, 041	5. 977	7.569
10	HEMBA1003987	48.695	30.080	108. 473	25. 632	23.222	28.008	21. 302	24.940
,,	HEMBA1003989	47.841	51, 466	128. 889	32. 288	24. 298	24. 627	15. 392	23. 174
	HEMBA 1004000	36.424	35.098	34. 843	16. 292	19, 541	20.604	16. 803	21.872
	HEM8A1004006	8.411	42. 393	12.931	2.863	3. 395	0.000	4. 943	9.742
	HEMBA 1004007	135. 300	114.014	286,000	90. 971	64. 473	74. 153	71. 985	79.319
	HEMBA1004010	58. 331	152.845	38.786	18. 676	18, 819	35. 229	31, 514	80.599
15	HEMBA1004011	62.306	16. 294	38. 336	12. 356	13.756	29. 583	26. 091	7.986
,0	HEMBA1004012	47.010	38. 053	139.110	42.415	22. 159	34. 340	27. 215	32.550
	HEMBA1004015	24, 416	26. 249	27. 372	12. 243	13.962	25. 082	25. 133	12.259
	HEMBA1004024	149, 457	114.788	479.037	80.579	77.896	75.066	57. 366	93.859
	HEMBA1004029	81.485	31.944	43.520	19.897	20. 191	38.768	36. 482	19.376
	HEMBA1004038	26.629	15. 823	19.708	12. 109	7.832	14, 400	12.855	17.771
20	HEMBA1004042	8, 177	10.678	12.830	6.612	11.484	7.963	11. 320	10.405
	HEMBA1004045	24.675	30.855	37.128	20.069	23.538	15. 509	17. 299	17. 447
	HEMBA1004048	95. 795	48. 977	78.760	36.608	40.779	45. 132	47. 334 23. 320	63.844
	HEMBA1004049	55. 947	543. 954	47.428	49.034	19. 297	56. 209	30. 423	68.865 32.553
	HEMBA1004051	69.776	31.608	51.948	13.046	25.684 15.007	38. 632 23. 414	13. 218	23.973
	HEMBA1004053	29.222	70.670	84, 481 34, 928	8.151	5, 353	28, 619	15. 237	14.807
25	HEMBA1004055	39.564 136.121	23. 202 122. 072	413.353	75. 363	81.883	66. 439	41.004	85.794
	HEMBA1004056		11.826	29.995	9.507	4.910	13.895	8.679	8. 388
	HEMBA 1004060 HEMBA 1004061	17.642	13.460	20.009	16.913	8. 228	14. 145	12.424	5.810
	HEMBA 1004067	165.029	79. 589	104. 390	62.419	50.783	89.115	94.004	91.850
	HEMBA1004071	28. 405	34, 722	37.707	19.775	14.692	17.342	23.864	27.554
	HEMBA1004074	128.445	51. 388	148. 050	35.606	37.851	50. 216	53. 461	46.373
30	HEMBA1004078	26. 126	14.714	20.940	9.721	16.211	17. 398	17. 388	14.057
	HEMBA1004085	42.006	24. 067	36.862	15.417	17.609	19.555	28. 362	21.993
	HEMBA1004086	27. 330	49, 843	21.238	43.213	24.232	16.260	12.409	22.262
	HEMBA1004097	45.296	15, 292	27.795	13.971	26.928	26.002	33. 192	19.361
	HEMBA1004100	40.930	37.210	48. 942	23.245	10.184	25.744	21.452	28.594
	HEMBA1004103	101.036	101.281	184.668	64, 176	44. 322	55. 385	41.050	40.000
35	HEMBA1004110	89, 903	65. 107	57.751	43.841	27.836	21.315	27.631	34. 280
	HEMBA1004111	171.907	134.108	296.310	95.474	115.874	78. 450	80.011	98.760
	HEMBA1004124	177.408	71.838	103.065	37.865	46.198	68. 531	109. 364	77.083
	HEMBA1004130	64.543	54. 797	171.602	50.628	35. 382	25, 601	19. 599	23.097
	HEMBA1004131	41.654	24. 184	33, 975	26. 913	23. 365	28.790	20. 022	24. 999
	HENBA1004132	55, 906	42.840	162. 243	42.708	30, 251	28.863	19, 780	22. 237
40	HEMBA1004133	64.624	30, 838	38. 522	29.390	20.897	28. 027	28.747	33.333
	HEMBA1004138	61.197	21.853	23.858	17.376	9.337	30.080	17.345	22.082
	HEM8A1004143	15.715	9.656	21.209	10.565	10.539	14.067	11.441	9.994
	HEMBA1004146	40.893	21.789	90. 537	30,633	32.870	23. 542	14.368	20.982
	HEMBA1004148	59, 990	18.796	22.167	11.049	17. 531	18.309	29. 374	22.628
45	HEMBA1004149	16.284	11. 131	18.385	7.758	7.634	7.577	5.890	13.683
45	HEMBA1004150	5. 223	4. 403	4.468	3.044	2.553	2.158	2.062	2. 260
	HEMBA1004154	111.110	40. 836	69.965	31.437	46. 253	58. 472	62. 983	47.866
	HEMBA1004164	139.670	107. 565	315.189	77. 326	47.327	57.372	46.726	67. 257
	HEMBA1004168	24.042	16. 530	18.698	9.347	9.400	13.838	3.054	13.060
	HEMBA1004199	22.894	9. 047	10.461	8,631	7.704	7.849	6.889	7. 253
50	HEMBA1004200	33. 301	51. 362	83.462	26. 185	27.548	17.580		32, 109
-	HEMBA1004201	54.766	23. 783	32.370	17.449	21.835	22.123	25. 993	20.006
	HEMBA1004202	14. 526	10.484	12.784	6.804	5.704	9.594	8.672	11.673
	HEMBA1004203	47.655	20. 140			14.171	19. 946	16.079	18. 151
	HEMBA1004207	6.344	3, 206	T1. 421	3, 936	6.145	5.704	21.692	7.780
	HEMBA1004210	33.071	43, 543	33.120	16, 340	43.156	21.814	19.639	15.015
55	HEMBA1004225	73. 182	63, 749	226. 133	59.585		32.703	25. 781	40.078
	HEMBA1004227	83.820	31. 222	42.541	16. 931	17.786	28. 177	25. 468	30.978

	HEMBA1004235	99.954	57.144	62.536	27.672	34. 345	69.613	47, 182	38. 807
	HEMBA1004237	27.504	21.542	17.029	18. 289	11.697	19.212	12.031	15. 922
	HEMBA1004238	79.210	38. 454	102.493	34. 130	27.841	36.089	27, 438	34. 578
5	HEMBA1004241	5. 863	2.654	7. 035	2.556	1.072	2, 912	4. 422	1.294
	HEMBA1004242	256.862	65.757	191, 327	80.010	76.455	85. 478	89, 242	62.567
	HEMBA1004243	72.699	55. 276	60.764	28. 287	47.148	36, 800	28. 491	47.743
	HEMBA1004246	44. 915	30. 967	100.300	22.414	17. 109	15. 470	12.686	18.700
		66.750	16. 238	24.674	18. 889	22.763	31.897	38, 415	17.377
	HEMBA1004247 HEMBA1004248	13.953	18. 412	17. 581	11.953	11.378	14, 538	12.794	9.562
10		24, 439	10. 494	10, 631	5.401	5. 142	14.218	12, 652	11.965
	HEMBA1004250 HEMBA1004252	37.349	20.650	22. 246	9. 949	9. 550	14, 570	21.841	18, 200
		10.994	19, 320	16. 415	15. 707	20.374	13, 845	11, 265	19. 838
	HEMBA1004260	22.716	14.715	13. 358	7. 615	5. 234	12. 282	15,089	11.397
	HEMBA1004264	235.310	195.750	654. 331	171.071	174. 292	115.073	102.973	144. 125
	HEMBA1004267	28.776	19. 025	23. 678	13.063	12.012	15. 529	14. 123	14. 593
15	HEMBA1004272	62.157	50, 491	53, 598	30. 356	36.472	42.005	58, 020	51.617
	HEMBA1004274 HEMBA1004275	70.423	38. 514	45. 176	17. 443	18.132	34.031	36. 295	22. 171
	HEMBA1004276	33.630	4. 481	14.011	9. 548	9.099	14.035	10.406	8. 615
	HEMBA1004279	16.536	11.082	13.356	14.834	7. 333	10.255	8, 919	12.068
	HEMBA1004284	29.688	30. 297	64. 483	13.658	17.646	17.327	17.630	13.770
	HEMBA1004286	32. 471	16. 566	18.049	12. 391	6.773	17.625	23, 811	13.547
20	HEMBA1004289	81.573	62.930	165, 571	49.704	34.785	37, 379	28. 939	41.740
	HEMBA1004293	72.466	34. 902	48.669	32.705	17. 408	57.764	53, 695	45.065 -
	HEMBA1004295	37.595	12.116	29, 975	11.634	5.514	25.018	23. 797	20. 926
	HEMBA1004302	10.880	5. 912	7.885	10.025	5, 190	6,060	5. 264	9. 355
	HEMBA1004306	426.811	177. 321	335. 168	107.645	123. 947	256.397	251.772	134. 005
0.5	HEMBA1004312	37.953	30.864	105, 533	30.747	25.847	16.140	16. 283	24. 272
25	HEMBA1004314	29.396	23. 332	95. 584	22. 179	18,544	11.015	8.804	20.974
	HEMBA1004321	47.670	29. 150	105. 316	35.655	23. 139	31.309	29.736	47.858
	HEMBA1004323	87.295	65. 931	221.440	44.690	41.425	36,609	34. 117	39.135
	HEMBA1004327	65.869	21. 284	21. 540	11. 985	14.419	27. 213	27.030	20.118
	HEMBA1004329	67.920	44.687	132.755	32.977	21.556	32.356	17.478	26.773
30	HEMBA1004330	8.765	7. 655	16.827	7. 164	3.843	9.511	7.660	4.615
30	HEMBA1004334	16.438	21.355	31.680	15. 109_	26.670	13, 368	10. 581	13. 568
	HEMBA1004335	204. 961	102.859	125. 226	69. 979	64, 392	78.772	71.641 89.558	83. 525 40. 224
	HEMBA1004341	186.677	30, 208	61.439	15. 995	41.404	87.221 26.797	42.054	59.071
	HEMBA1004344	261.676	76, 316	123. 332	42.705	51.432 21.953	33.115	33.526	36.846
	HEMBA1004347	65.249	32.610	97. 858 29. 441	37.038 19.803	18.786	23. 126	19, 103	18.719
35	HEMBA1004349	22.353	35. 727 65. 544	237. 050	49. 039	34. 141	32.597	28.166	46.343
	HEMBA1004352 HEMBA1004353	75.508 54.322	66. 042	132.169	40.563	27. 380	39. 551	30.556	56.886
	HEMBA1004354	43.687	29. 352	79. 264	22.784	20, 533	21.755	16.860	22.429
	HEMBA1004356	44.730	22. 201	27. 487	10.404	8.280	22.159	16.039	15.038
	HEMBA1004360	91,412	28. 429	71.634	26.232	36.259	59. 602	38. 361	50.410
	HEMBA1004365	9.956	10.099	14. 263	5. 481	5.631	5.802	6.791	6.167
40	HEMBA1004372	3, 613	4, 593	5. 338	0.000	1.638	1.507	3. 555	1.568
	HEMBA1004377	53.834	41.410	47.048	29.140	26, 163	34. 545	30.827	33. 572
	HEMBA1004389	20.540	22.800	24.474	14.497	13.968	16.620	14. 951	17.114
	HEMBA1004391	60.284	22, 653	44.013	14. 283	19.018	31.716	23. 931	23.617
	HEMBA1004393	177.786	197.548	108, 554	32.455	75. 399	76.587	39.777	44.665
	HEMBA1004394	28. 949	11.849	12. 442	5.544	10.440	17. 825	10, 981	8. 836
45	HEMBA1004396	37.907	26.956	102.760	18.571	16, 519	15.025	13.681	21.980
	HEMBA1004401	22.519	21.858	30.601	14.945	13. 592	15.418	20. 530	20.774
	HEMBA 1004405	42.933	38, 835	117. 844	34. 528	23.557	19.155	18. 506	29.842
	HEMBA1004408	50, 497	27. 151	55, 000	25.559	15. 351	19. 522	15. 546	20.863
	HEMBA 1004414	45.769	51.722	64, 316	19.655	19.324	39.735	26. 527	36.385
50	HEMBA 1004429	61.867	59.067	190.058	39.014	50.304	38.462	27. 517	46.317
JU	HEMBA1004433	49, 568	39.828	146. 938	37. 521	28. 383	24.241	24.651	42.005
	HEMBA1004440	31.849	22.499	37. 132	18.742	22.366	23. 183	21.969	35. 073 37. 635
	HEMBA1004444	59.488	46.586	163.763	30.695	29.990	24.833	10.855	16.078
	HEMBA1004446	22, 134	12.309	29. 426	11.920	3.385	14.862	21.159	27.156
	HEMBA1004451	31.688	21. 251	28. 136		15.678	19.695 17.680	14, 173	7.972
55	HEMBA1004452	36.593	5. 268	18. 479		5. 737	24. 494	22.897	29.042
	HENBA1004454	50.056	27.897	32.786	24.382	20.631	1 (4. 434	1 4.031	1 23.042

					9 : 304 1	71 700 1	74 444 1	46 444 1	EE 045)
	HENBA1004460	138. 550	96.143	356.058	74.883	54.735	70.698	38. 344	55. 945
	HEMBA1004461	64.074	19.163	24. 808	12.846	16.373	23, 508	22.827	10. 137
	HEMBA1004468	134. 439	72.774	210.409	77. 409	60.142	55.229	42. 361	49. 457
5	HEMBA1004479	82.994	32.899	92. 282	35.519	44.435	43.183	31.836	37.116
	HEMBA1004482	5.602	7.682	11.248	36, 034	2.925	5. 535	5.693	5.972
	HEMBA1004491	16.736	6. 285	17.615	14.018	11.729	19.804	10.683	11.280
			71.456	148. 355	58. 479	48, 596	46, 968	46.648	57.279
	HEMBA1004499	94. 095			7. 282	9. 979	2.762	3. 174	13. 389
	HEMBA1004502	21. 523	9.344	18. 265				22.873	15.812
10	HEMBA1004505	26.042	15. 980	43. 855	17. 516	15, 469	22. 190		
10	HEMBA1004508	12.004	29. 395	42.664	21.849	21.426	14. 469	9. 224	9.845
	HEMBA1004507	96.377	87.688		103. 472	34.160	81.068	54. 939	151.142
	HEMBA1004509	52.657	14. 880	19. 120	15. 228	17.009	24.783	24. 565	13.476
	HEMBA1004523	20, 156	18.209	11. 197	16.529	14,651	13.004	20. 267	19.467
	HEMBA1004528	42,620	27.819	48, 069	14. 426	25. 267	43.038	40. 239	37.718
	HEMBA1004534	75, 090	41.159	44. 399	31.300	16.686	31, 317	21.009	18.589
15	HEMBA1004536	31.531	13.343	23. 664	14, 085	4, 408	13, 133	12. 981	15. 201
			181.508	233.819	97.018	122.402	183. 507	100. 197	150.062
	HENBA1004538	352. 363				12.324	17. 317	22.764	15. 212
	HEMBA1004542	47. 360	17. 733	29. 238	17. 280	39.674	38, 686	23.830	33.542
	HEMBA1004552	63.401	29. 585,	26. 857	43. 567				
	HEMBA1004554	62.231	11.953	25. 084	16.273	32.820	33.809	20.812	5.536
20	HEMBA1004558	30.217	12.133	31.036	15.840	30.638	65. 183	19.155	30. 921
20	HEMBA1004560	68.901	16.566	17. 908	16.431	10.034	30.093	22. 379	22.683
	HEMBA1004564	48.119	14. 911	35. 565	31.983	32.464	30.028	20.965	32.479
	HEMBA1004566	32, 479	29. 553	20. 970	32.788	42.949	40.715	23. 273	32.960
	HEMBA1004573	17.728	13.843	7.118	9. 972	19.952	9.755	9. 278	8.100
	HEMBA1004576	39.572	26.733	42.044	10.704	17.505	12.441	17. 232	25. 361
	HEMBA1004577	46.233	11.570	97. 881	39, 434	13.437	41.089	34.426	35.314
25	HEMBA1004586	82.532	71.398	213.814	70. 289	52.589	45.729	23.395	38. 312
	HEMBA1004596	72.534	32.493	45.820	27.585	27.854	34, 997	33.847	38.473
	HEMBA1004604	99.019	48. 582	103. 587	35.723	49, 392	48.377	56.558	69.256
	HEMBA1004607	53. 557	37.013	100. 999	27.559	26.143	28.796	21.692	42.044
	HEMBA1004610		14. 854	69. 908	15. 349	12.120	9.108	8.858	15.087
		20.690		42. 426	22.819	15. 568	10. 691	6. 697	10.317
30	HEMBA1004617	22.592	20. 386		49. 931	29, 836	29. 902	12.194	27. 438
**	HEMBA 1004622	78.025	46.803	209.059			17, 193	15. 579	20.821
	HEMBA 1004626	38. 170	36.312	110.684	22.791	14.118			
	HEMBA 1004629	33.858	37.886	87. 440	53, 228	47.341	28. 160	12.170	28.096
	HEMBA 1004631	35. 946	10.475	4. 434	7.390	17.128	22.775	9. 569	32.852
	HEMBA 1004632	27.084	13.891	23, 598	10.209	7.802	11.754	22.566	6.362
25	HEMBA1004633	78.391	33.135	114.054	17. 197	49.008	60.659	48.857	40.810
35	HEMBA1004636	52.397	20.706	34. 962	10.085	22.609	21.255	13. 502	25. D39
	HEMBA1004637	4. 228	4.304	6.747	5.278	9.756	4.086	2.597	5.024
	HEMBA1004638	0, 241	0.000	0.000	1.008	0.000	0.000	0.113	0.000
	HEMBA 1004645	57.971	29. 263	111.067	32. 545	17.998	27.214	20.560	24.845
	HEMBA 1004656	16, 139	9.194	21, 399	12.766	18.216	14.099	17.122	12.004
	HEMBA1004657	20.820	23.742	69. 842	9.422	138. 932	42.697	9.048	13.383
40	HEMBA1004666	7. 321	3.174	18.097	5, 962	9.830	5.098	2. 525	7.512
	HEMBA1004669	94. 910	36.291	111.210	30.591	20.021	28.018	25.500	25.624
	HEMBA1004670	57.231	17.070	60. 538	23.280	13.173	24. 312	23.413	14. 342
	HEMBA1004672	63.471	50.154	145.619	39.883	31.559	25.617	20.328	28.099
	HEMBA1004689	152, 993	93. 435	103.311	81.212	50.901	83.998	57.329	84.276
	HEMBA1004690	28.240	10. 247	13. 401	8, 159	4. 952	13.963	13.991	11.785
45	HEMBA1004693		15. 228	20. 803	14, 290	13.070	16.726	9.014	13.531
		18, 359			58.849	34.416	51.983	42.641	50.271
	HEMBA1004697	81.532	48.847	148. 587		42.224	58.936	64.906	37.506
	HEMBA1004702	97.518	62.966	49.904	20.714		38.377	24. 626	31.783
	HEMBA1004704	99.561	48.717	236. 687	38.866	33.457			
	HEMBA1004705	12.717	12.313	40, 950	9. 649	17.803	10.638	5.969	4.810
50	HEMBA1004706	33. 516	9. 825	16. 175	10,779	10.830	17. 906	13.036	12.703
	HEMBA1004709	51. 126	39. 934	136, 723	32. 285	25.072	21.674	15. 230	23. 755
	HEMBA1004711	46.766	9. 203	57,020	12.805	14. 304	16.154	12.982	9.790
	HEMBA1004723	121.283	47.643	73. 497	30.236	56.917	55.719	56.298	52.009
	HEMBA1004725	56.905	32.051	70.171	12.221	48.208	34.021	35.739	12.501
	HEMBA1004730	36.072	10.037	30.016	7.633	13.361	7. 545	8. 989	34. 832
	HEMBA1004733	30.769	29. 884	23.348	6.988	2.998	8.055	8.031	2.822
55	HEMBA1004734	11.912	11.974	36. 595	3.988	12.556	7.653	4.303	15.670
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	HEMBA1004736	55.309	25, 331	132.333	45, 653	38, 696	23.516	19.970	34, 509
			20, 004	172.069	29. 562	22. 161	14.904	12.665	18. 349
	HEMBA1004748	53.832			27.788	33, 184	60.214	44.636	42, 250
5	HEMBA1004749	127.285	45, 137	73.698					43.505
·	HEMBA1004751	81.283	64, 830	173.888	54. 165	36. 368	41.802	31. 283	
	HEMBA1004752	59.058	32. 785	109.428	32.254	29.090	34. 259	30.970	33.029
	HEMBA1004753	204.044	247. 466		156.689	102. 755	131.323	83.515	263.882
	HEMBA1004755	57.638	59.677	83.850	22.148	29. 800	30.642	13.064	23. 261
	HEMBA1004756	9.965	16. 228	11.023	8. 349	5.780	9.109	111.628	14. 885
40	HEMBA1004758	36, 487	26.558	116.970	22. 341	14. 553	14.773	11.840	14. 406
10	HEMBA1004763	67.343	19.641	33.742	13, 841	16, 720	25.489	23.061	18. 650
	HEMBA1004768	29.177	24.043	38. 303	6. 573	10, 298	3. 197	10.352	13. 391
	HEMBA1004770	10.327	14. 492	10.901	6.416	6.310	7.963	10.868	7.955
		46.910	34, 314	76.491	31,609	22.830	23. 102	30. 433	32. 358
	HEMBA1004771		28. 705	63. 968	24. 931	18.754	43.049	32.720	26.795
	HEMBA1004775	39.253		10.103	5.466	9, 000	16.400	10.105	8. 046
15	HEMBA1004776	22.604	11.017				32.337	24.057	43. 529
	HEMBA1004778	78.144	77.681	223.475	37. 540	33. 791		9.479	6. 329
	HEMBA1004784	9,826	18. 370	102.812	8.313	15, 151	11.373		
	HEMBA1004785	25.723	16. 345	26. 216	6.651	10.649	10.674	13.732	11.615
	HEMBA1004789	18.173	14.508	16.096	7.804	8, 691	10.011	7.713	11.389
	HEMBA1004795	14. 283	12.973	25.122	11.028	9, 351	9.757	9. 905	12.028
20	HEMBA1004797	65.927	33.745	73.888	34, 142	28. 246	40.067	32.715	25. 583
20	HEMBA1004803	36.634	41.124	65.880	27.072	30.957	22.607	22.520	26.554
	HEMBA1004806	11.997	8. 183	21.467	8.868	9.653	9,000	7.894	8:199 -
	HEMBA1004807	16.352	14.481	22.459	11. 249	12.009	13.340	7.935	9.118
	HEMBA1004816	29.782	24.075	95. 884	18.110	29. 259	8.180	12.578	10, 934
	HEMBA1004820	8.636	7.466	8.862	4. 249	4.018	4.269	6.876	3. 493
05	HEMBA1004833	159.947	50.729	81.248	38.650	64.754	83.155	56.657	65, 121
25	HEMBA1004847	51, 456	25. 570	40.694	21.115	36. 572	35.053	31.537	40.529
	HEMBA1004850	77. 254	24.014	38.620	21.854	26, 080	54.413	50. 197	24.185
	HEMBA1004853	57, 117	32.704	72.480	23.951	31.887	25.058	20.050	20.982
	HEMBA1004864	46.043	27.344	59.824	26.750	13.898	16.719	20. 308	17.843
	HEMBA1004865	12.257	14.642	31.748	44.090	14. 331	13.454	13.835	15.797
	HEMBA 1004880	56.788	50.021	126.837	35. 420	26.589	24.064	20.647	23. 264
30	HEMBA1004882	42, 450	18. 453	29. 340	16.782	13.013	13.652	10.676	19.977
	HEMBA 1004885	8. 545	4. 947	5. 350	4.891	2.933	3.711	3. 652	6.615
	HEMBA1004889	28, 103	22. 485	32.049	17.078	14. 363	23.391	15.605	16.916
	HEMBA 1004900	19. 922	15, 709	33.254	10, 423	9.045	6.539	5. 245	9.440
	HEMBA 1004909	88. 522	49, 269	163.284	48.147	35. 537	36.045	18.861	27.933
	HEMBA 1004918	64.384	43.134	105.868	34, 899	22. 323	24.073	15.857	25.370
35	HEMBA1004923	47.731	37. 996	69, 168	19.659	26. 441	18.192	10.213	20.111
	HEMBA1004929	11.048	14, 003	10.808	12.050	7, 539	9.882	8. 967	11.809
	HEMBA1004930	101.277	92.425	279.652	80.664	66.618	34.331	31.091	41.874
	HEMBA1004933	9.145	5. 566	12.895	7.786	12. 296	10.327	95.467	5.417
	HEMBA1004934	7, 311	7.106	43. 966	10. 208	4, 750	5.866	9.143	12.805
	HEMBA1004937	43.331	27. 219	38.802	15. 368	17.734	15, 280	15.784	46.365
40	HEMBA 1004943	51.072	26.833	32.001	21.614	16, 458	27.585	29. 628	38.533
	HEMBA1004944	84. 363	46.788	126. 294	41.803	28. 989	38.514	31.589	23.074
	HEMBA1004946	64.638	28.144	37.908	17.163	24. 332	27.854	34.636	31.712
	HEMBA1004952	90.835	18. 893	40.862	12.824	20.090	33, 568	20.062	19.020
	HEMBA1004954	14.656	36, 003	41.485	27.126	23.696	20.777	6. 946	29. 261
	HEMBA1004956	5. 975	9, 923	6.635	7.743	0.953	4.578	1.565	5.188
45	HEMBA1004950	86.030	77. 420	136.061	60.735	49. 221	47.560	29. 646	45.929
,0	HEMBA1004971	31.046	5, 439	7. 559	12.468	17, 946	16.068	19.705	18.480
	HEMBA1004972	77.318	38. 259	56. 654	35.819	27. 295	40.233	30.004	50.710
		35. 524	13.502	16.731	9.641	11.726	14.716	19. 197	22. 580
	HEMBA1004973	6.756	9. 870	11.419	9.684	29. 373	8.701	2.217	10.523
	HEMBA 1004977		11.088	13. 909		5, 158	5.699	2.642	10.106
50	HEMBA1004978	8. 589	33. 440	87. 268			16.453	11.605	22.124
-	HEMBA1004980	34.091		17. 944	9, 205		11.553	6.083	5, 456
	HEMBA1004982	14.750	8. 271 13. 488	20. 831	11.831	3. 348	10.309		8. 305
	HEMBA1004983	38. 285		26. 297	18. 434		22, 491	24. 452	33.683
	HEMBA 1004995	27. 256	28.515	33. 238		7.816	13. 134		14. 698
	HENBA 1005004	13.855	10.490	22. 502	18.478		28.617	18. 581	16. 940
55	HEMBA1005008	64.714		19.462			26. 432		23.116
55	HEMBA1005009	34, 543	15.673	1 13.402	10.043	1 19. 122	1 .4. 742	1 .2. 333	
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Table 32

					A0 404 T	40 144 1	44 644 1	88 777	22 016
	HEMBA1005019	49. 250	24. 872	25. 349	20.834	30.144	32.629	20.777	27.016
	HEMBA1005021	37.224	63.713	38.065	17.061	15. 647	20. 369	25. 859	32.656
_	HEMBA1005029	30. 265	17.783	35. 352	16. 531	19.588	26.517	15. 798	16.604
5	HENBA1005035	393. 404	200. 167	574.746	133.872	210.689	177.872	156. 563	170. 510
	HEMBA1005036	115. 345	41.961	73.015	39. 541	44. 451	66.623	55.833	51.349
	HEMBA1005039	28.850	19.922	57.018	13.971	11.999	26.427	15. 206	7.350
	HEMBA1005047	93.995	31.868	54. 335	18. 576	28.338	31.562	31,930	23. 751
	HEMBA1005050	78.015	41,690	73, 330	29.830	26. 504	35.887	21.640	35. 653
				29. 553	15.707	7.836	15.618	19. 435	13. 336
10	HEMBA1005062	23.050	15.803				10.541	5.005	10.849
	HEMBA1005066	10.980	11, 364	31.553	13.509	5. 668			
	HEMBA1005067	39. 308	34. 578	39.795	44.519	24.643	21.272	19. 379	20. 121
	HEMBA1005070	73. 155	34, 949	68, 556	29.956	38.004	38. 211	48.007	31.733
	HEMBA 1005075	88.089	37. 798	148.675	40.537	33. 271	33.074	28.661	30. 201
	HEMBA 1005078	100.064	37.746	66,827	33. 115	41.170	55. 560	51. 231	17, 112
45	HEMBA 1005079	137.757	86, 238	294, 118	73.304	76.035	75.084	47. 255	76.170
15	HEMBA 1005083	18. 102	7.642	17.087	6.711	6.184	8. 675	9. 287	9.609
	HEMBA1005084	82.712	38. 248	47.063	26.664	27.435	37. 552	38.419	28. 349
	HEMBA1005088	31.610	22. 435	76.774	22.700	18. 926	23.875	8.895	12.447
	HEMBA 1005089	68. 944	55. 156,	178. 226	34.742	32.350	38. 545	22.869	28. 148
	HEMBA1005090		86, 156	117. 997	94.811	57.034	81.098	54. 187	116.066
		148.861		63. 940	33.378	33.962	48. 589	35.467	36.021
20	HEMBA 1005096	83. 125	30. 911		10. 197	13.998	34. 420	22. 696	15. 975
	HEMBA 1005101	69.080	14.020	34. 136					21. 337
	HENBA1005107	82.659	25. 203	36.223	11.215	21.514	32.720	25. 972	
	HEMBA1005113	7.977	17. 225	31.501	7.563	44. 493	5. 157	6. 957	9. 761
	HENBA1005123	173.637	77. 260	555, 672	126.908	94.628	90.446	70.735	90.016
	HENBA1005133	58. 192	40.749	122.920	29.864	16.700	17.652	8. 802	18.988
25	HEMBA1005135	8. 259	9. 125	14. 962	2.213	16.732	6.892	3. 383	6. 189
23	HEMBA1005145	185. 299	101. 220	352.159	92.082	88.750	122.118	76.475	90.044
	HEMBA1005149	220. 122	109. 352	274.492	120.663	125. 192	96.704	92.083	128.030
	HEMBA1005152	125. 948	96. 291	226.882	58. 505	33.738	46. 323	27. 534	34. 457
	HEMBA 1005159	15. 760	11.274	9.399	6.198	6.191	6.861	12.001	4, 556
	HEMBA 1005172	1653. 208	89.658	73.666	54.667	33.118	55.680	32.520	70.907
	HEMBA 1005185	9.954	17. 248	10.492	34.452	3.558	3.117	6.026	11, 173
30	HEMBA1005186	23.745	10.048	27.091	13,067	7.719	15.412	15.086	15. 591
	HEMBA 1005195	14. 573	8.648	11.038	19.306	6.313	25.313	13.510	9. 183
	HENBA 1005201	52. 322	13. 197	47.505	13.091	12.078	8. 531	23.532	9.848
	HEMBA1005202	98. 566	30.141	71.588	27. 954	44. 381	66.294	42, 390	39.695
	HEMBA 1005204	184. 429	287. 155	382.039	168.753	203. 458	222.970	143.609	358.646
		148. 946	61.309	84. 791	34. 139	49.115	66. 295	83.608	76.159
35	HEMBA1005206				8. 236	8. 038	13.031	7.751	11.441
	HEMBA1005219	21.685	17.755	9.606	40. 547	32.099	53.017	26. 025	32.004
	HEMBA1005223	79.969	42.665					4.681	7.079
	HEMBA1005229	26.819	9. 926	21.841	3.135	5. 090	6.656		
	HEMBA1005230	71.184	67.313	201.065	79. 279	59.679	77. 484	47. 808	86, 511
	HEMBA1005232	7.374	6.386	17.522	8. 552	3.285	12.098	4. 975	3.965
40	HEMBA 1005238	96.780	44, 134	51.932	8. 128	20.776	69. 291	49. 474	35.019
40	HEMBA 1005241	142. 598	104. 185	428.635	18.773	78.033	74.434	42. 333	63.097
	HENBA1005244	76.771	32. 597	37.797	16.459	12.489	35.934	31.814	35, 602
	HEMBA1005248	241. 316	60.348	73.077	25.067	41.351	117.666	88. 193	54.014
	HEMBA1005251	37. 505	33. 247	108, 631	23.585	14.915	23.393	14, 302	16, 409
	HEMBA 1005252	53. 401	25. 532	37. 199	15.002	20.744	31.279	24. 207	27.562
	HEMBA1005267	17.238	39, 564	20.097	27.506	31.874	11.013	14. 526	14.024
45	HEMBA1005274	16.538	8.744	18.308	9.021	10.103	11.943	8. 914	11.978
	HEMBA1005275	69.133	43.329	216.468	46. 290	57.647	37.411	25. 040	41.913
	HEMBA1005288	65.401	50.495	150.714	33.833	34.633	28. 241	24. 910	40. 164
	HEMBA 1005293	17, 403	9, 430	23. 201	4.467	3. 192	25. 620	6.775	8.771
	HEMBA1005296	223.097	811.623	894. 835	738.361	220. 523	698.319	418. 435	1376.785
	HEMBA1005301	36.708	16. 970	29.798	11.929	12.544	22. 221	35. 726	32.270
50	HEMBA 1005304	83.978	71.914	260. 016	50.686	36.101	36. 160	24. 896	47.838
				74.215	27.494	27.352	34. 920	21. 424	38.882
	HEMBA 1005305	44. 218	33.773		13.836	7. 908	8. 958	9. 090	10.440
	HEMBA1005311	33.034	20.140	48. 263					
	HEMBA1005313	11.165	36, 175	17.550	7.047	11.502	14. 209	63.072	9. 124
	HEMBA 1005314	6. 948	2. 955	22.604	6.336	3.812	5.787	4.069	4. 891
55	HEMBA 1005315	72.349	54.139	156.842	34.545	43.132	26.415	28. 942	28. 442
55	HEMBA1005317	20. 230	8. 451	17. 258	9.796	11.664	9. 976	9. 263	8.017
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